

METHODS AND APPARATI USING SINGLE
POLYMER ANALYSIS
Inventors: Zhao et al.
Serial No.: TBD
Docket No. C0989.70049US01

Express Mail Label No. EV 292458974US
Date of Deposit: February 5, 2004
Docket No. C0989.70049US01

Figure 1

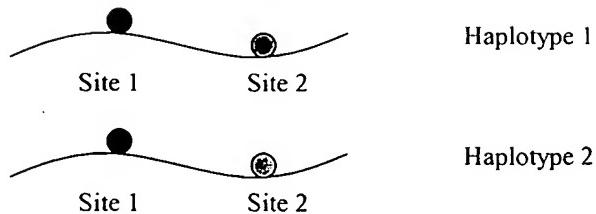
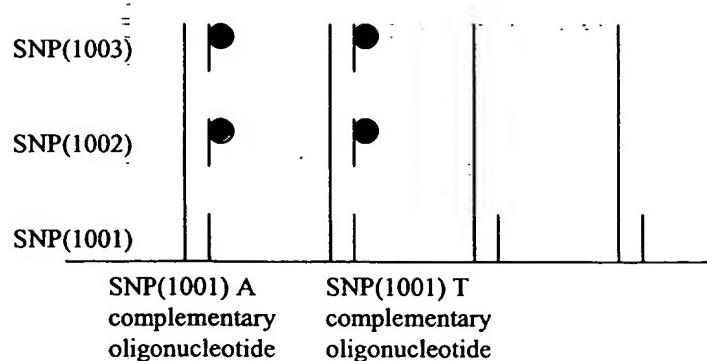


Figure 2



Figure 3



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Figure 4

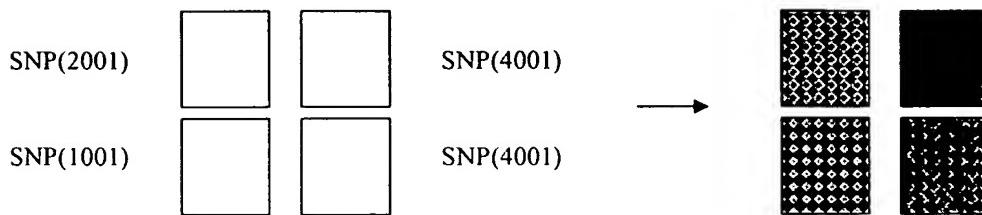


Figure 5

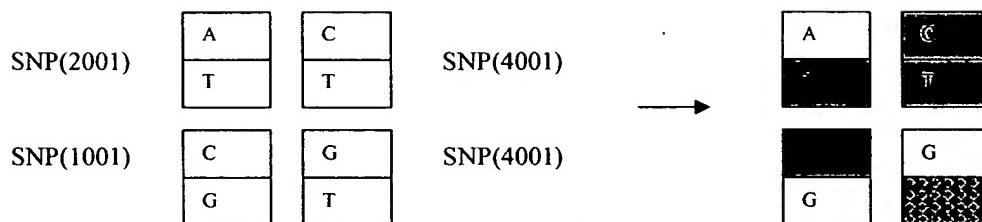


Figure 6

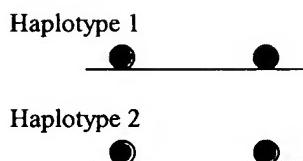
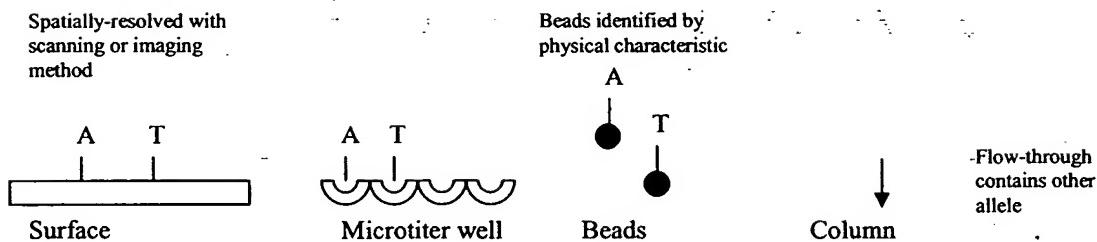


Figure 7



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Figure 8

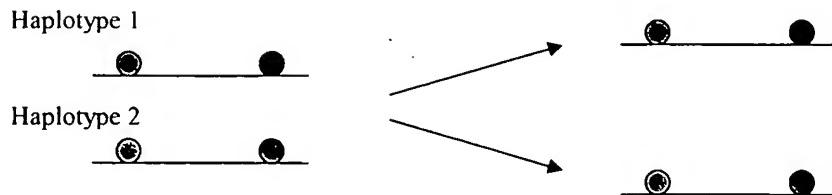


Figure 9

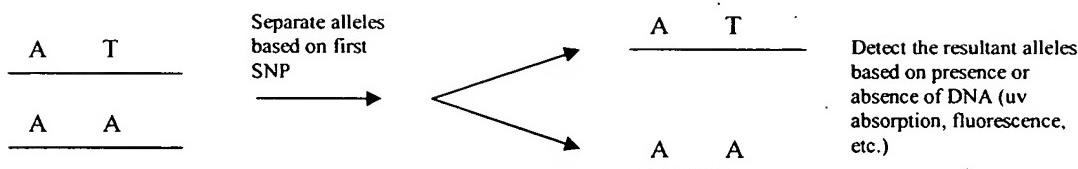


Figure 10

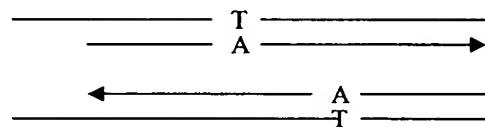
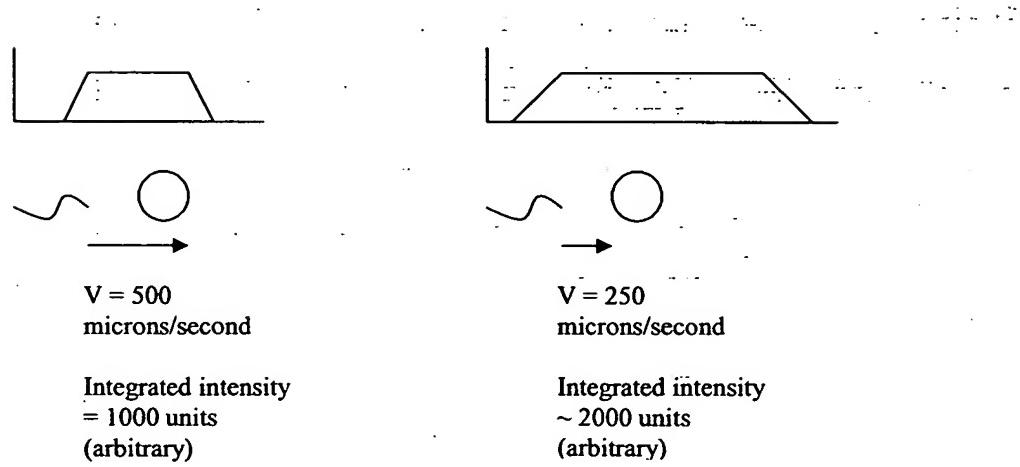


Figure 11



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Figure 12

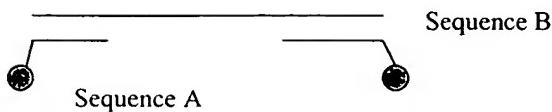
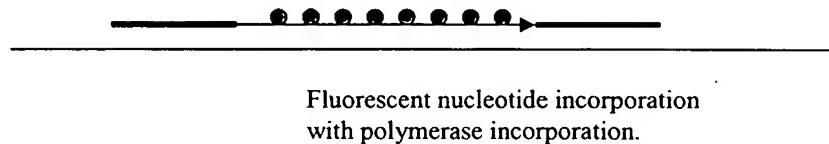


Figure 13



Fluorescent nucleotide incorporation
with polymerase incorporation.

Figure 14

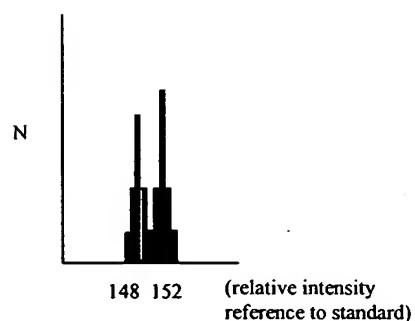
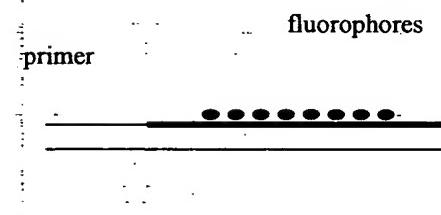
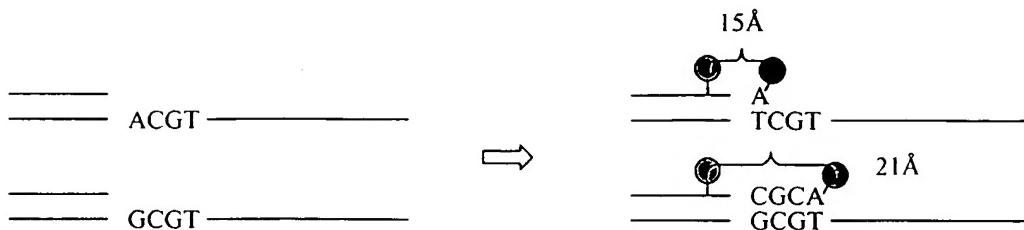


Figure 15



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Figure 16



A/G SNP detected through primer extension-spFRET and differential termination using ddATP

Figure 17

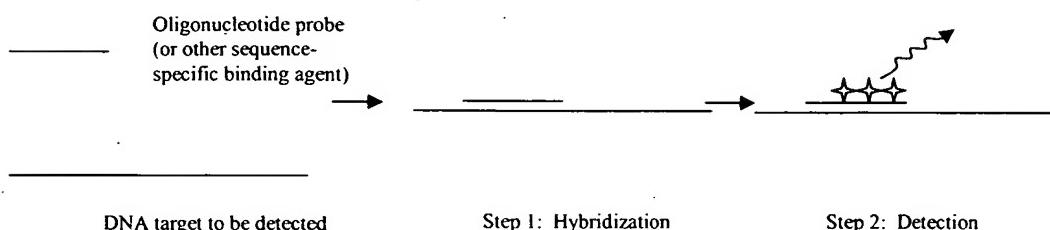


Figure 18

TWO-COLOR PRIMER EXTENSION ASSAY

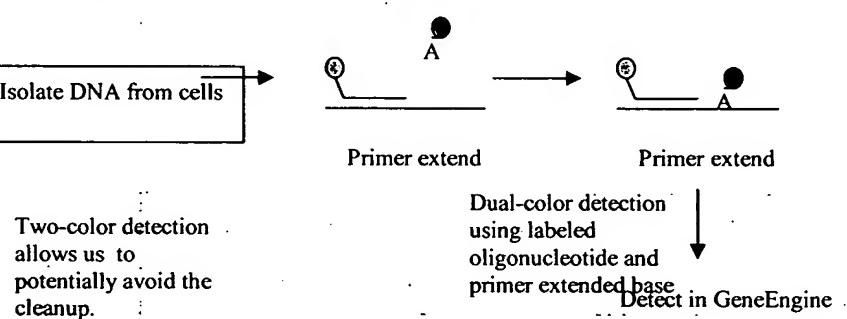
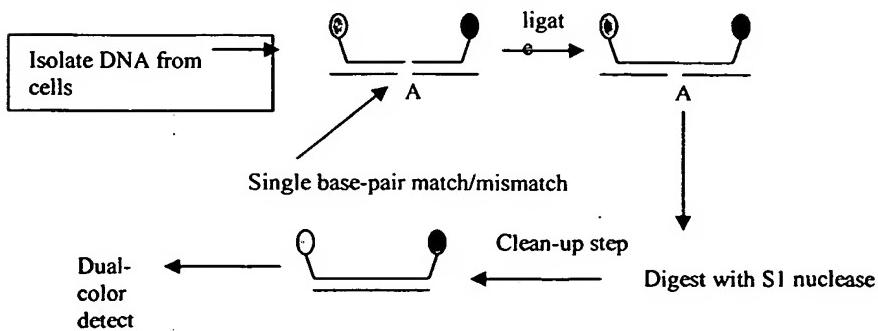


Figure 19

TWO-COLOR EXTENSION AND LIGATION ASSAY



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Figure 20

spFRET-BASED ASSAY OR PRIMER EXTENSION ASSAY-BASED CLEAVAGE OF PRODUCT (LIKE TAQMAN)

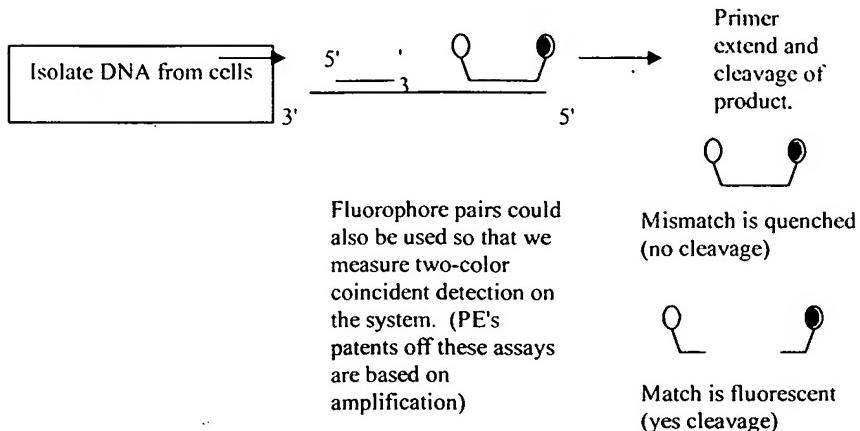


Figure 21

spFRET-BASED ASSAY BASED ON COINCIDENT HYBRIDIZATION

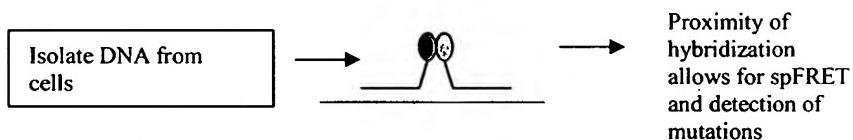


Figure 22

spFRET-BASED ASSAY WITH COMBINATION OF SINGLE-BASE EXTENSION REACTION

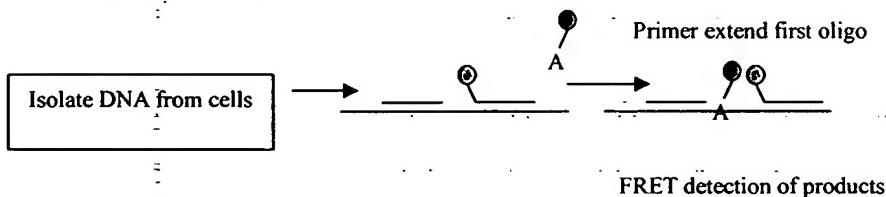
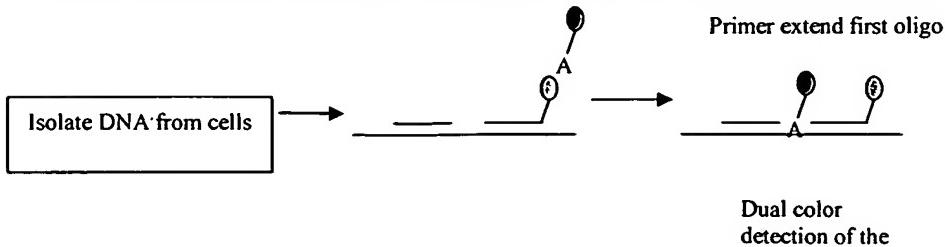


Figure 23

TWO-COLOR DETECTION ASSAY IN COMBINATION WITH PRIMER EXTENSION



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Figure 24

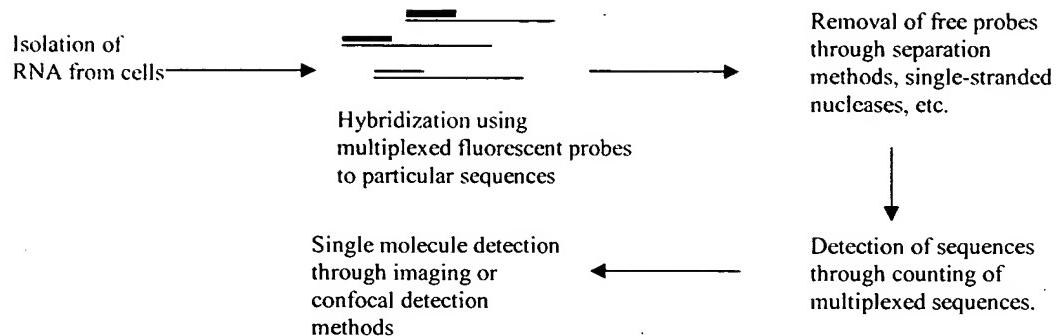


Figure 25

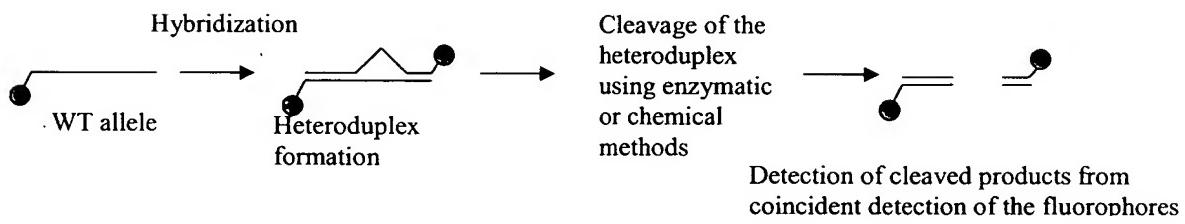
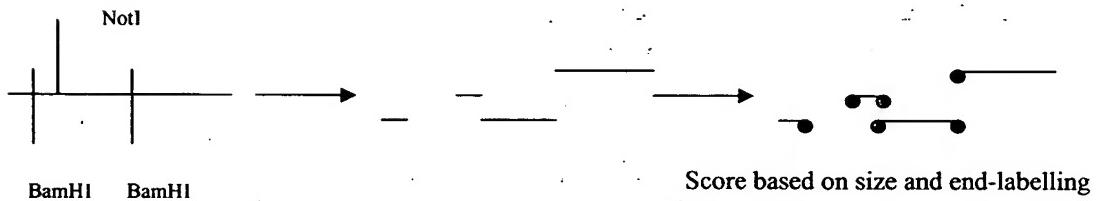


Figure 26



Single Molecule Fluorescent Tagging and Coincident Counting of Molecules

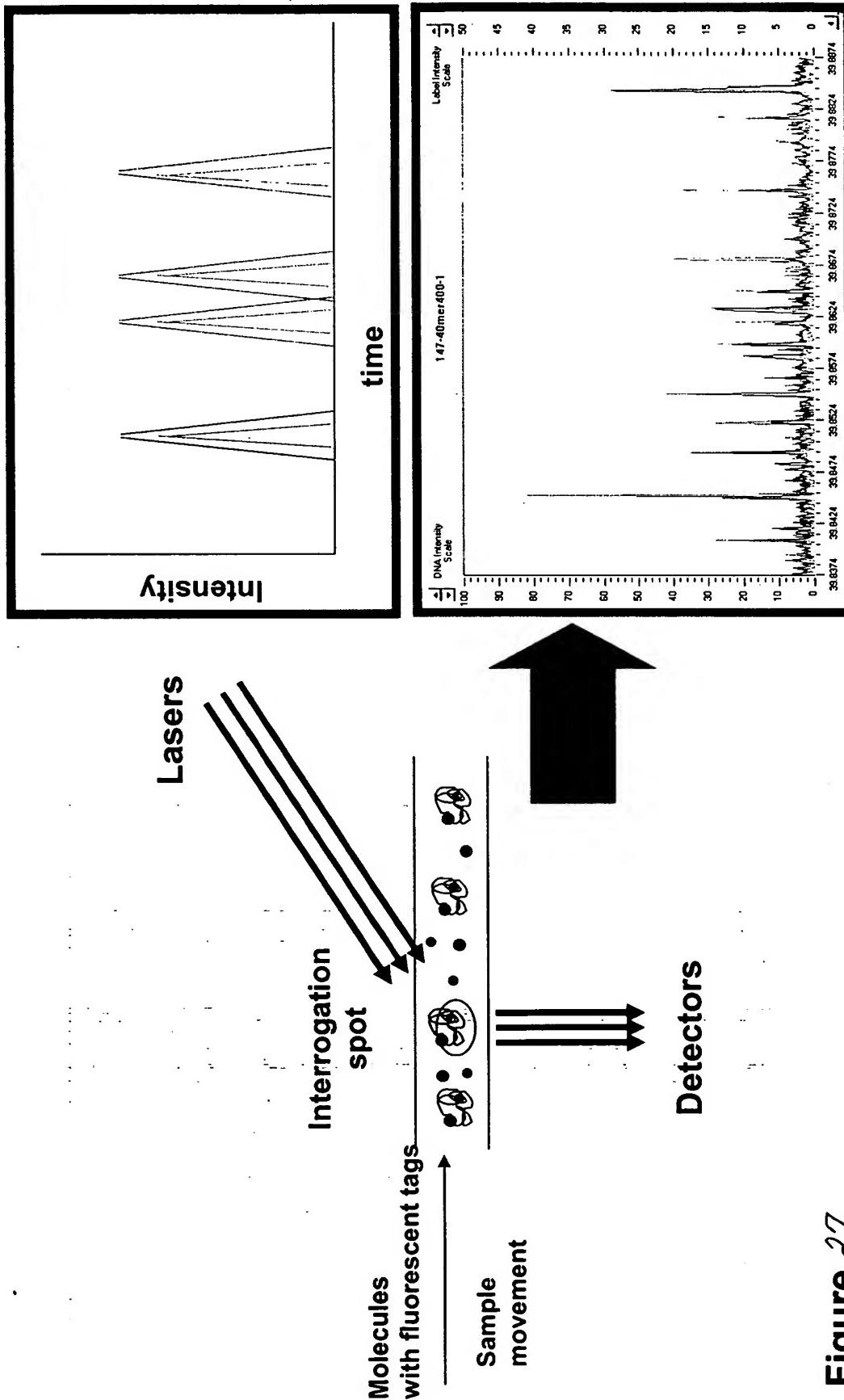


Figure 27.

Titration of Dual Labeled Oligonucleotide

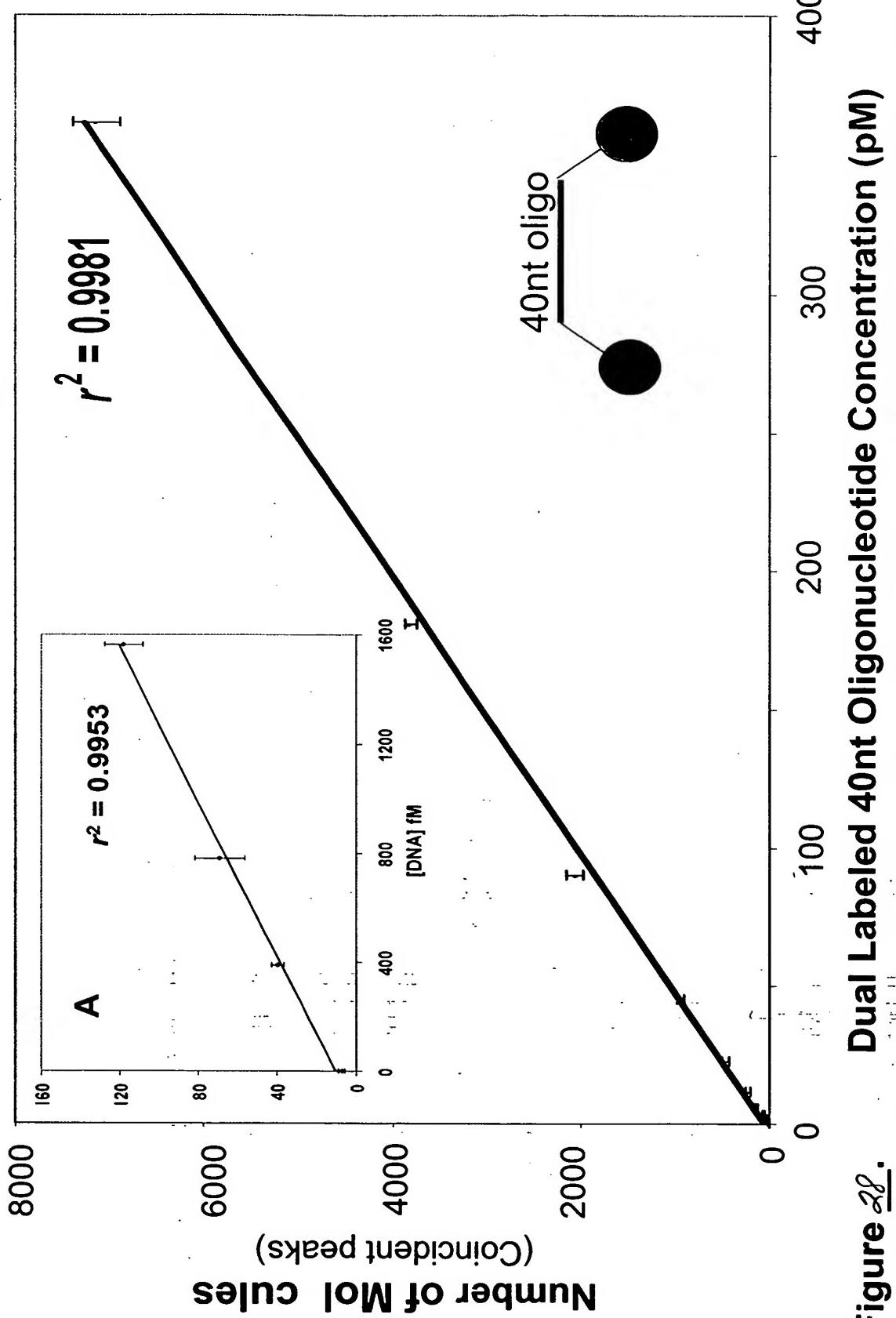


Figure 28.

Dual Labeled 40nt Oligonucleotide Concentration (pM)

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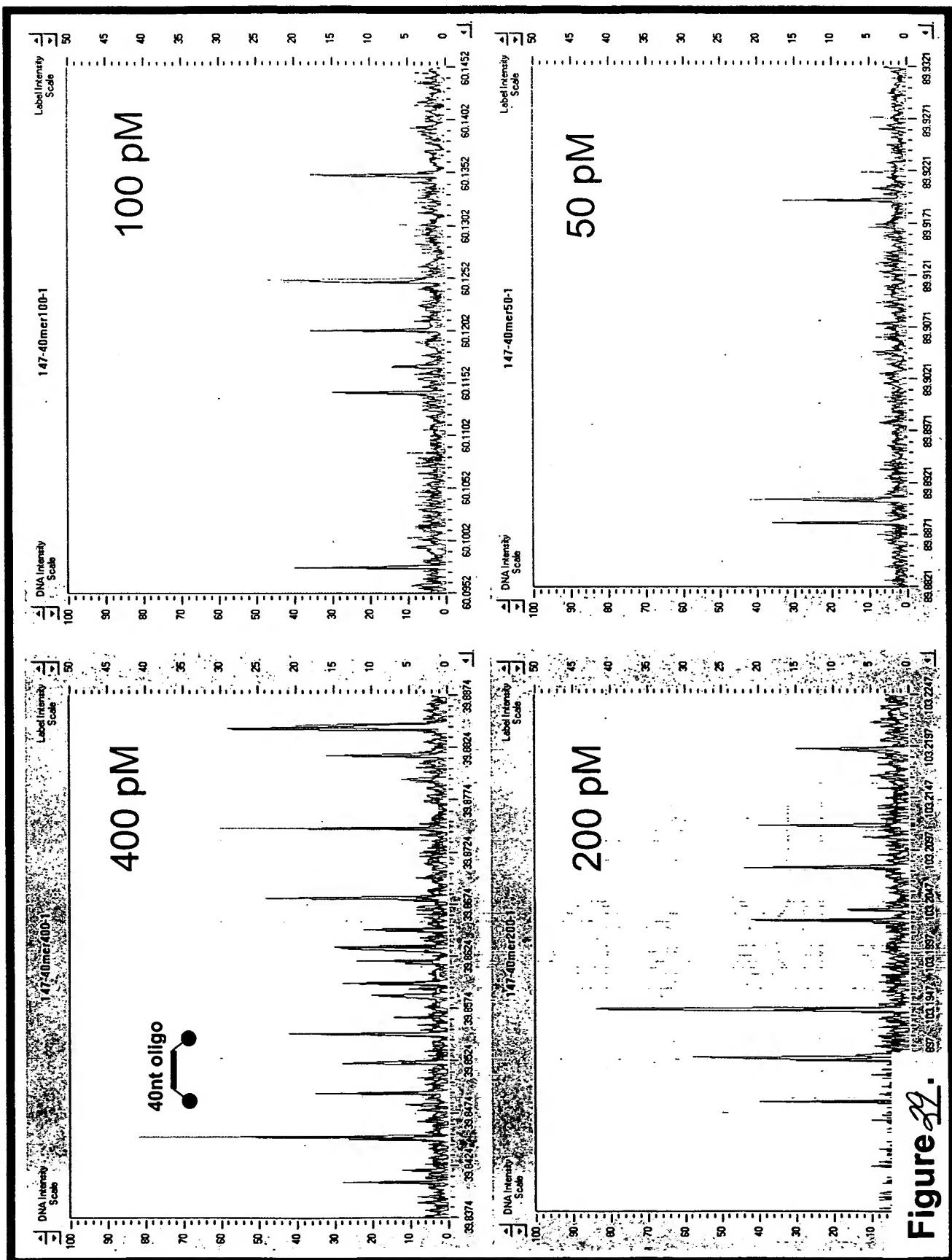
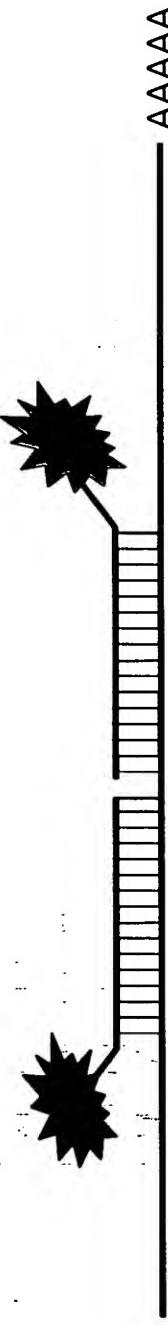


Figure 22.

Two Assays Validated

- **Dual Probe Hybridization:** Hybridize two oligos (20 - 30 nt) to RNA target



- **Probe Extension:** Hybridize one Cy5 labeled oligo and incorporate TAMRA by reverse transcription

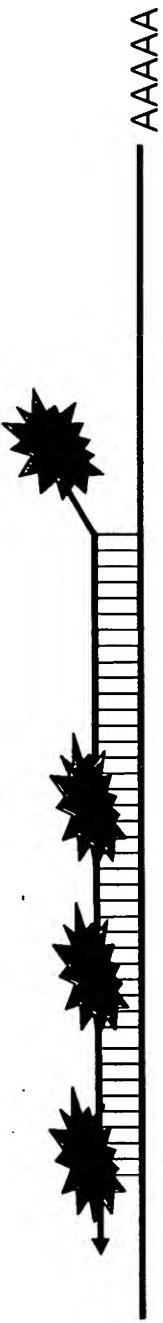
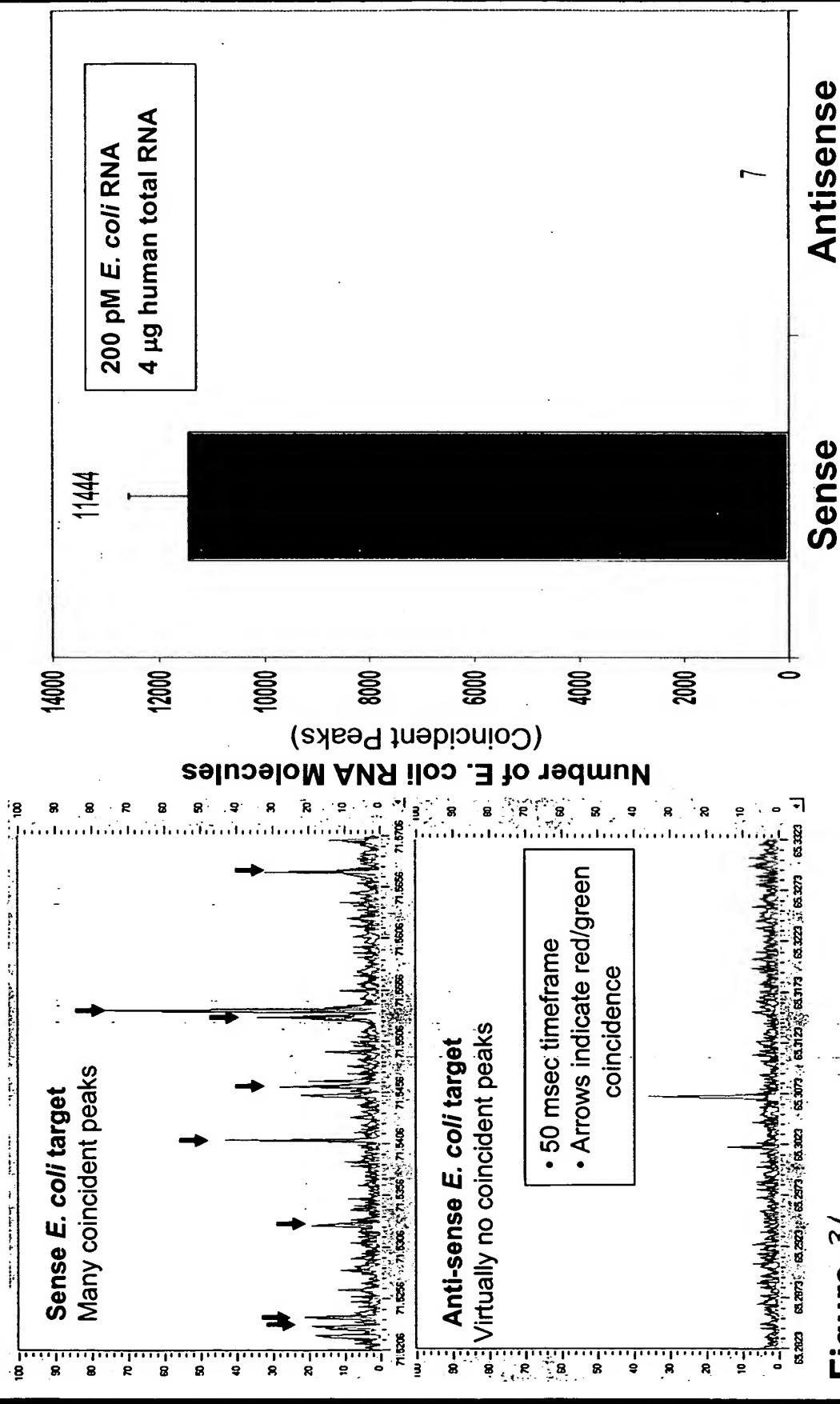


Figure 30.

Dual Probe Hybridization

- Sense vs. Antisense *E.coli* spike in total human RNA background



Probe Extension Assay

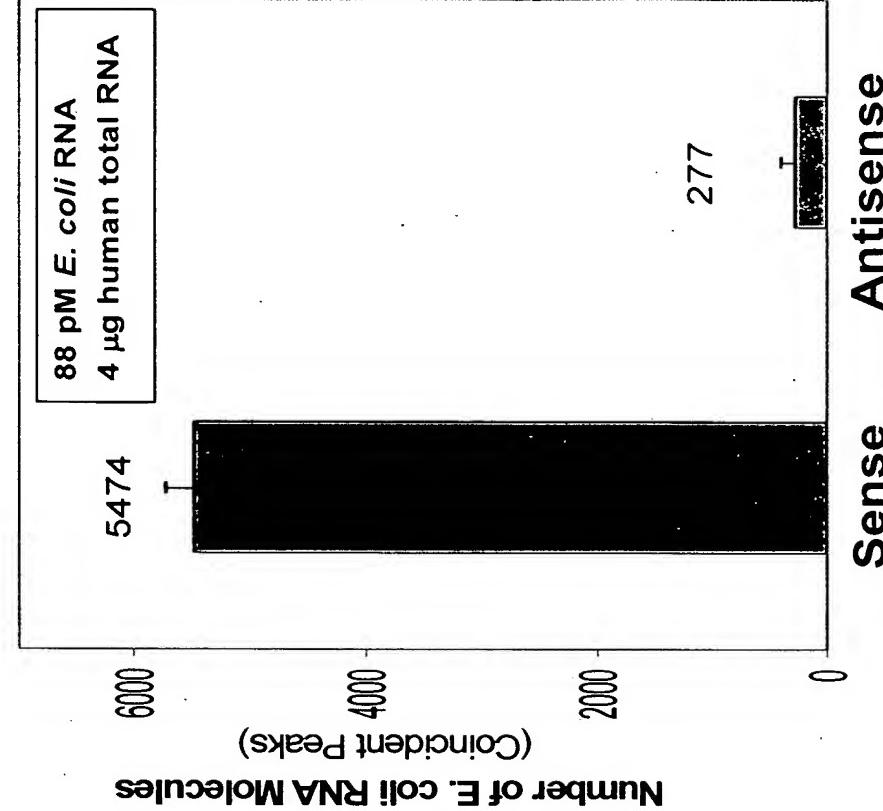
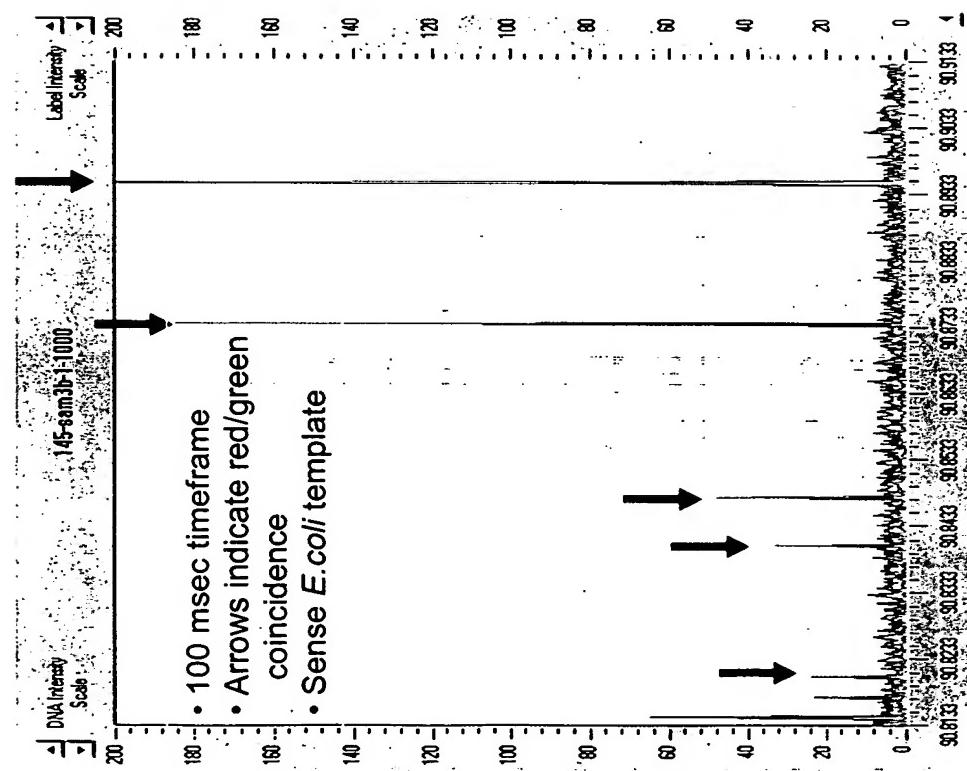
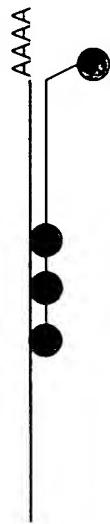


Figure 32.

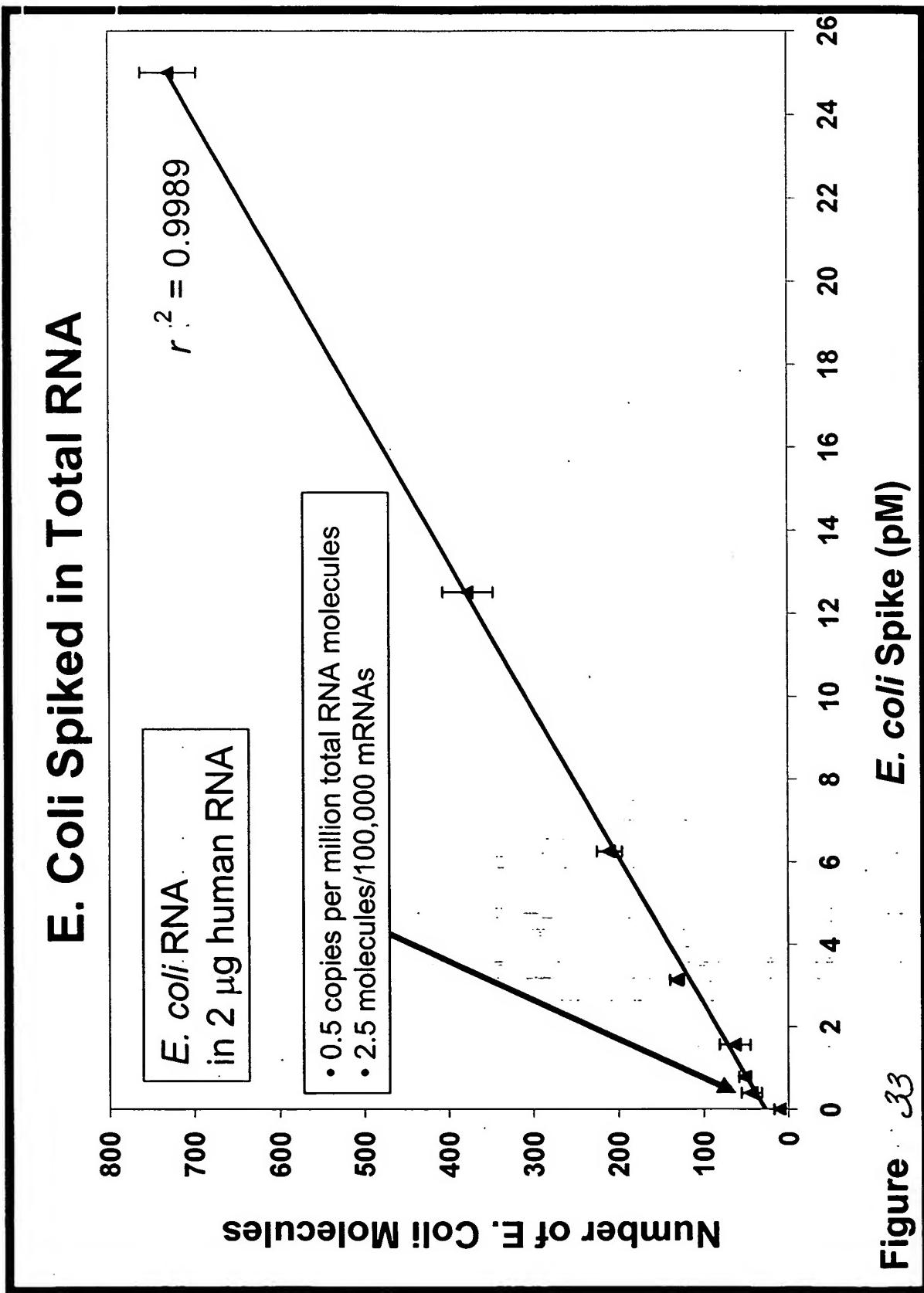


Figure 23

Lamin A/C and β -Actin Levels in Human RNA

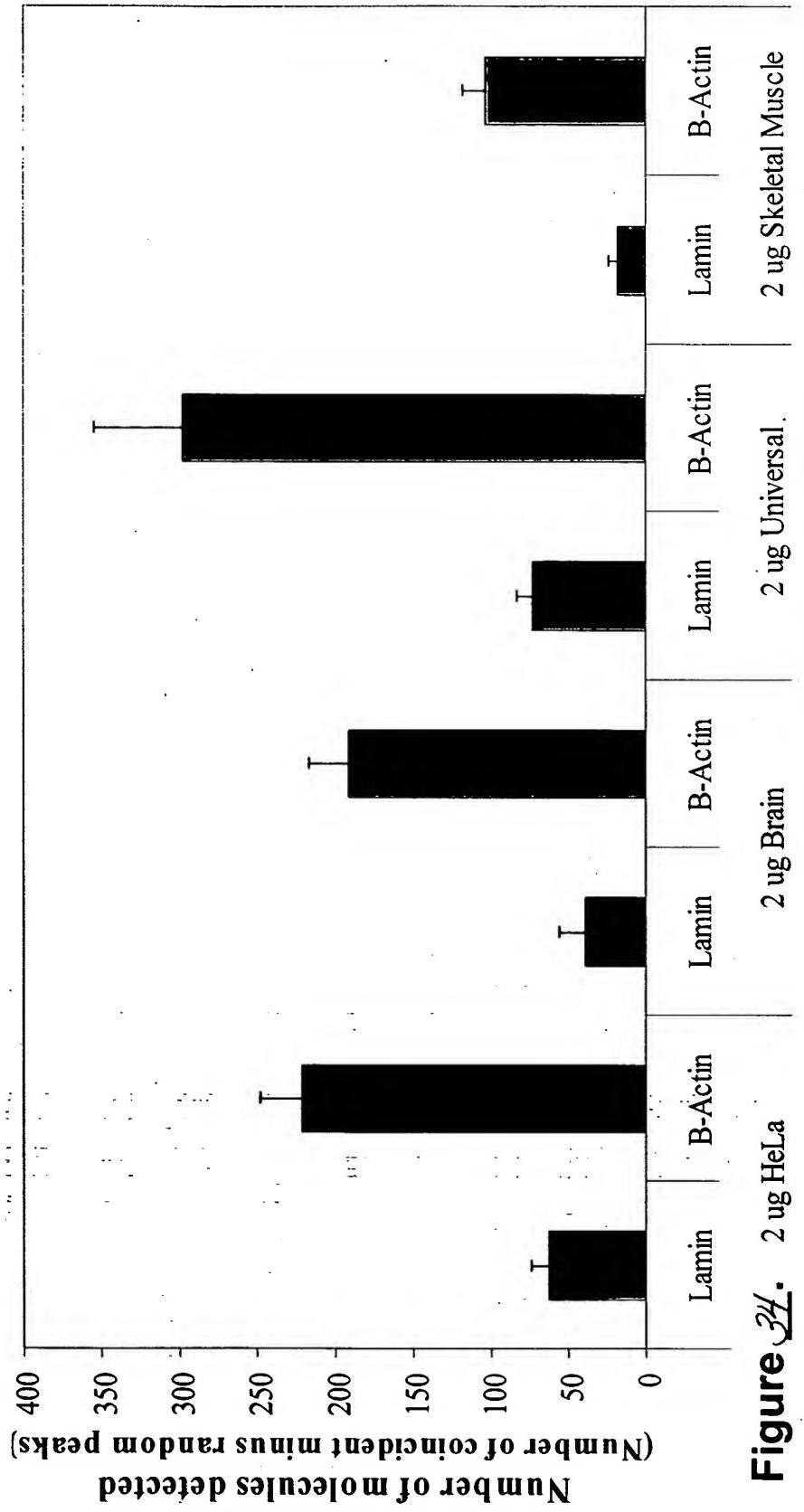


Figure 24. 2 ug HeLa

Quantitating Poly(A)⁺ Levels

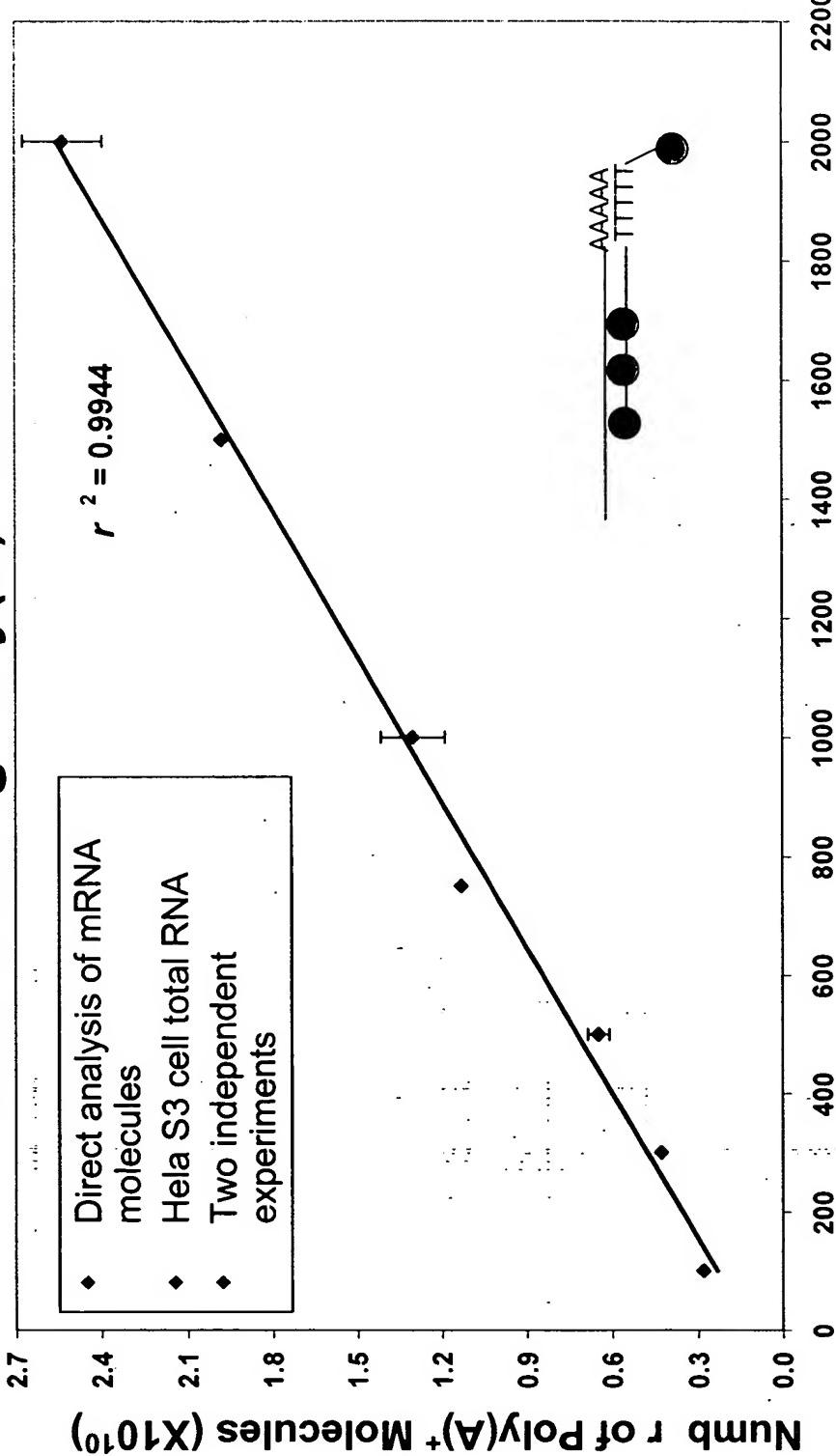
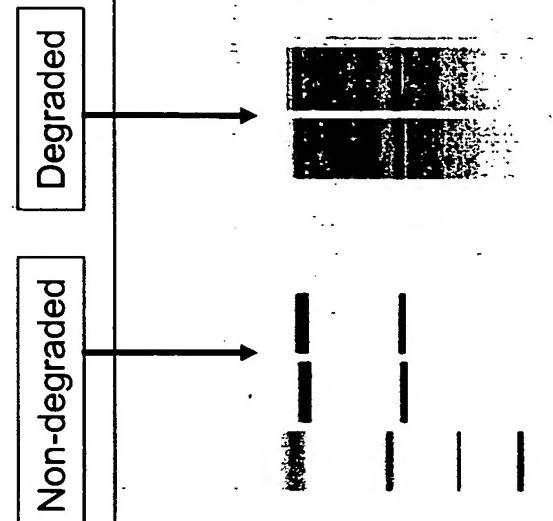


Figure 35.

Amount of Total RNA (ng)

Assessing mRNA Degradation

Total RNA Samples



Ratio of Green/Red Average Peak Areas by DirectRNA™

	G/R Avg. Peak Areas	Std Dev
Non-degraded RNA	2.64	0.02
Degraded RNA	0.88	0.10
Dual-labelled 40 mer	0.79	0.02

Figure 3C.

Gene X analyzed by *DirectRNA*™ and by Real Time-PCR

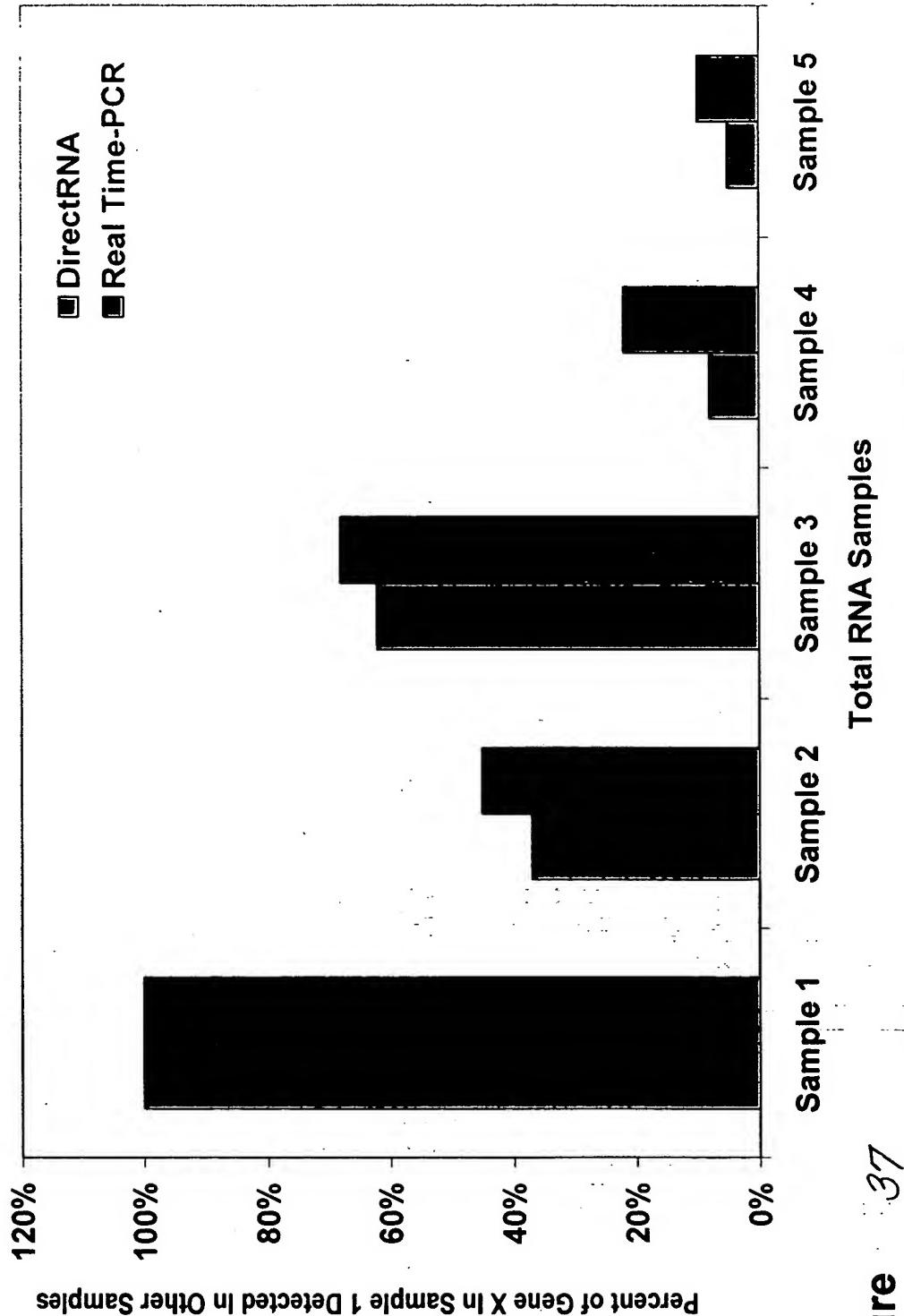
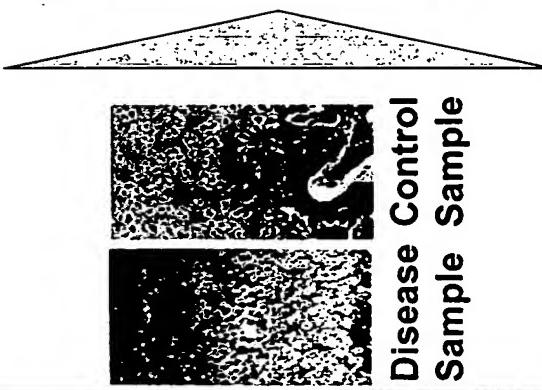


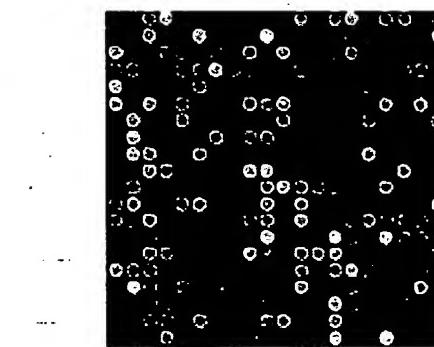
Figure 37

Quantitation in RNA Analysis by DirectRNA™

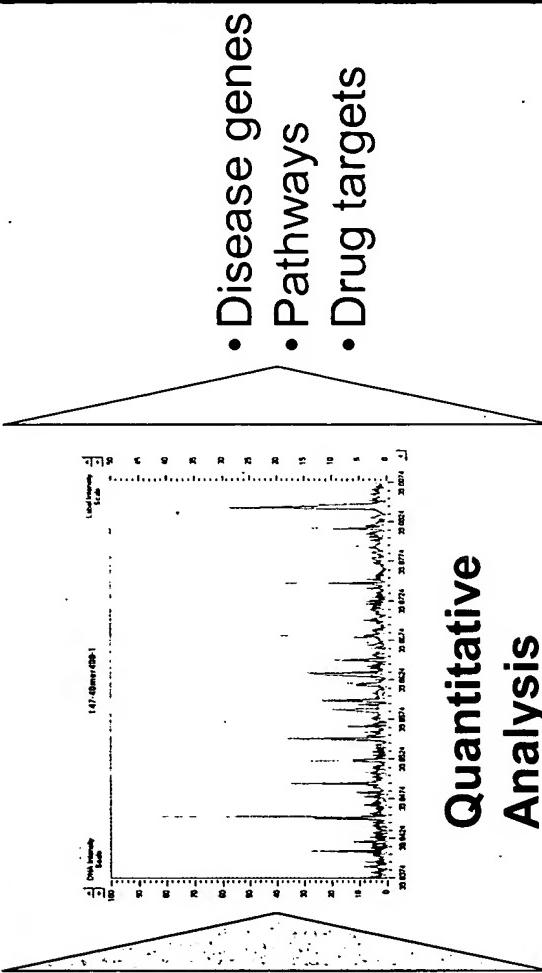
INPUT



TOOLS & ANALYSES



OUTPUT



First-pass analysis across thousands of genes

Detailed analysis of specific genes of interest

Figure 38.

Fig. 39A

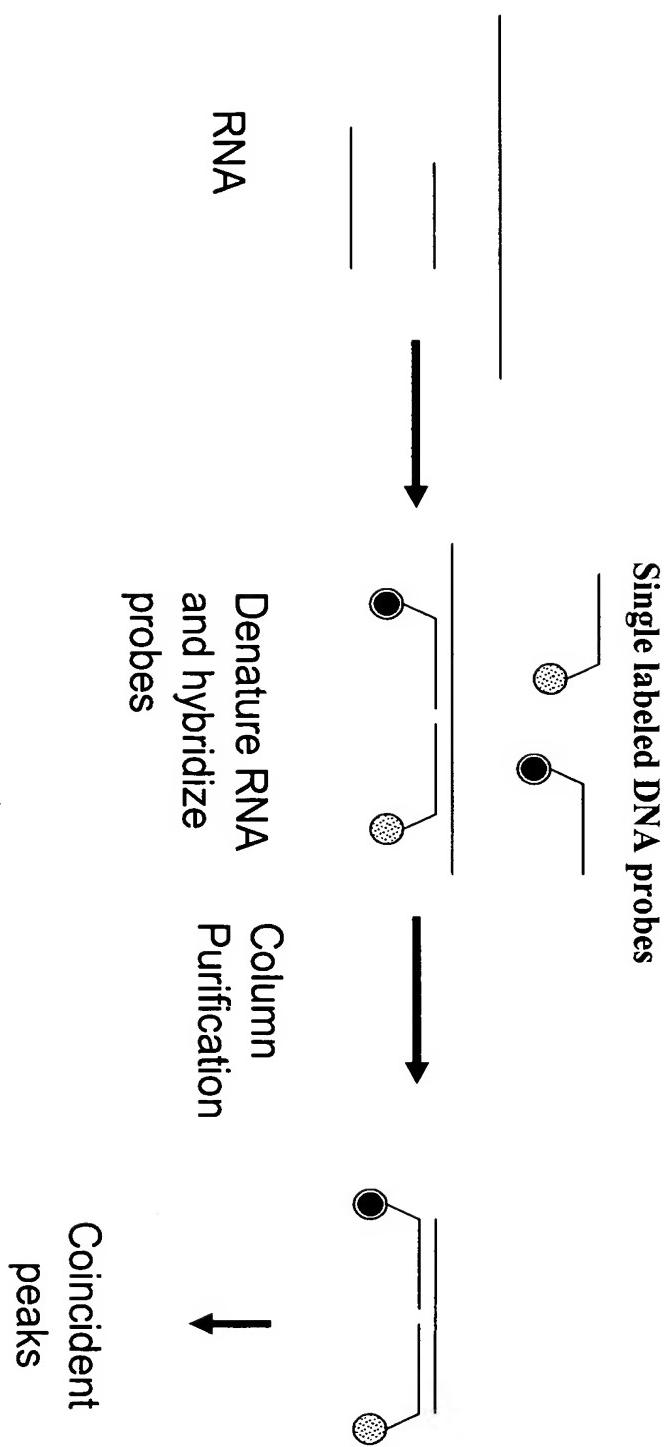


Fig. 39B

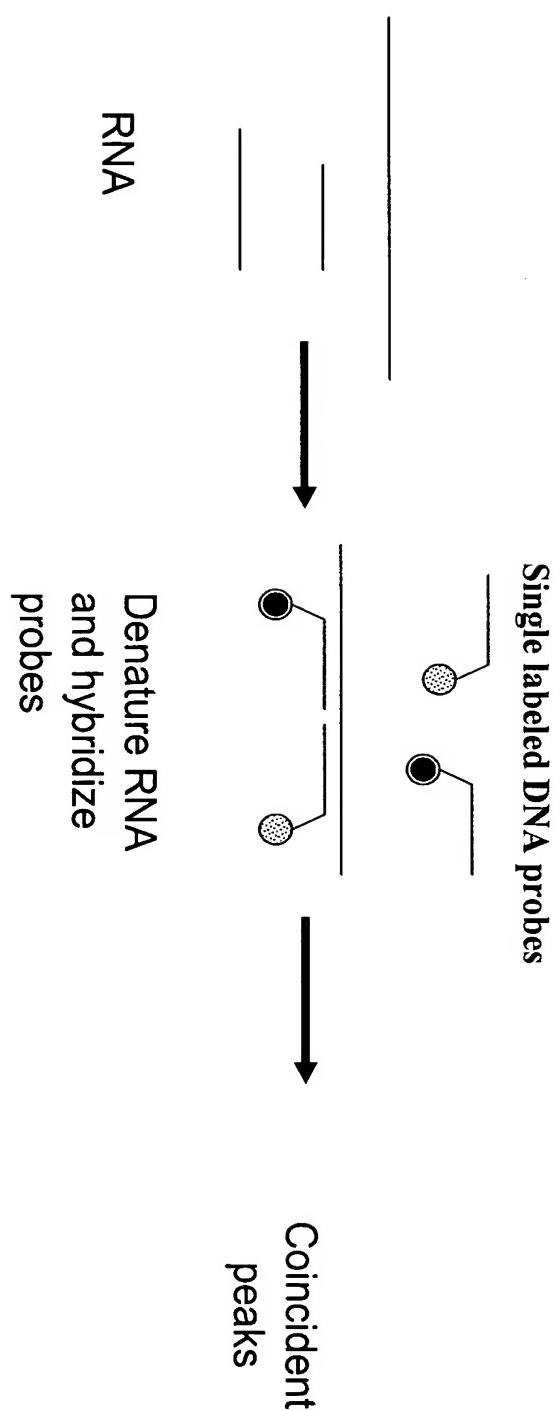


Fig. 40

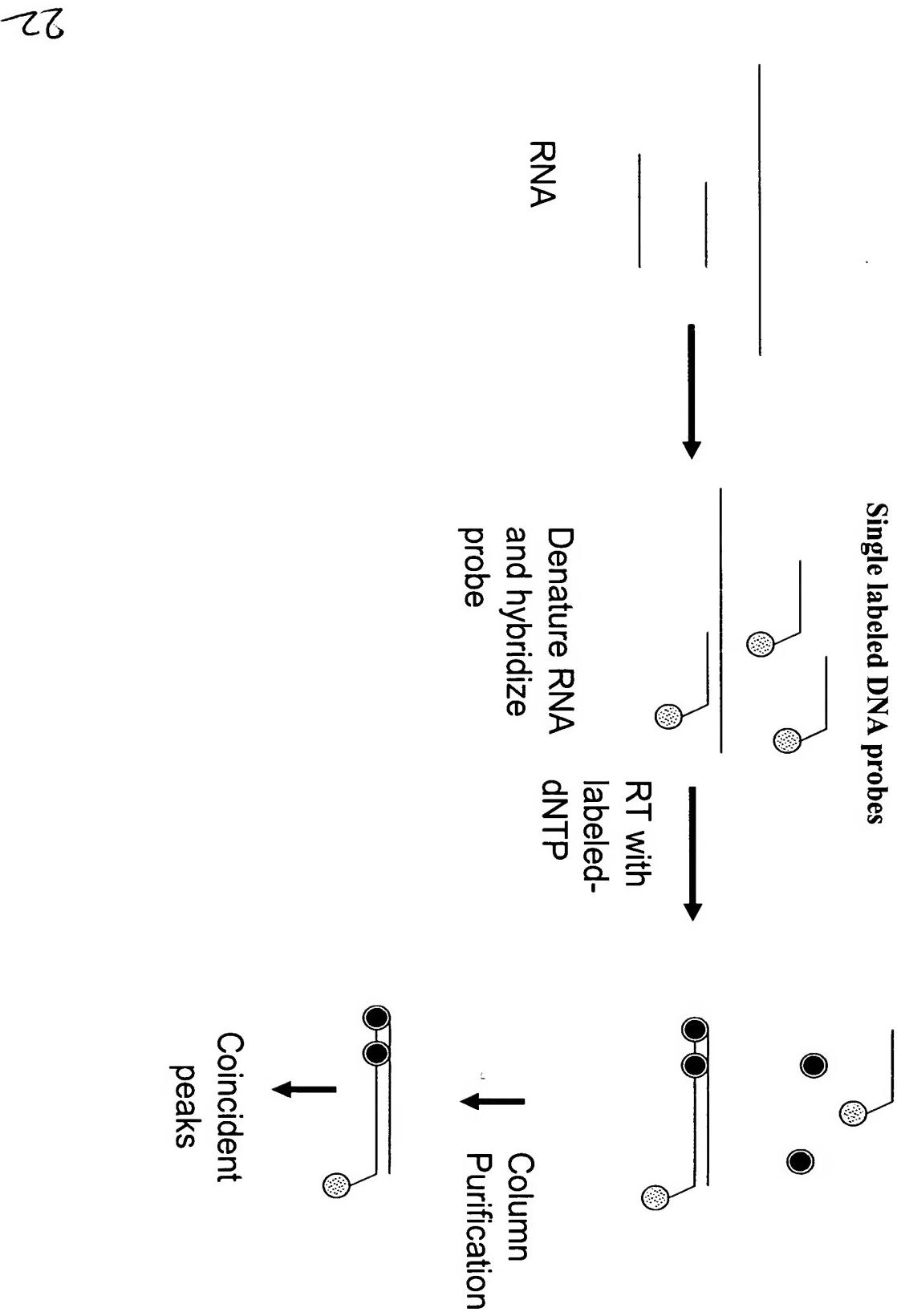


Fig. 4/A

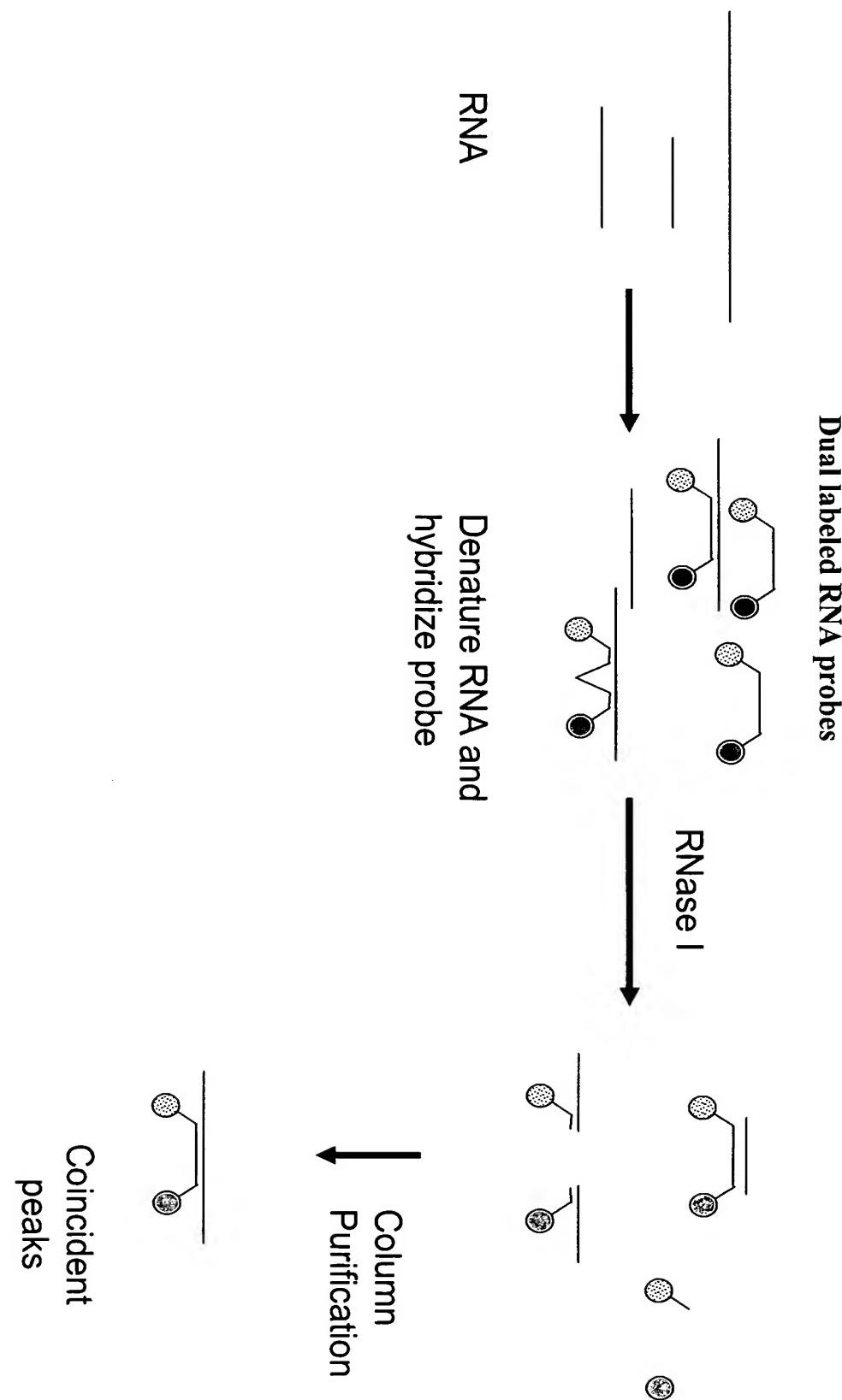


Fig. 4/B

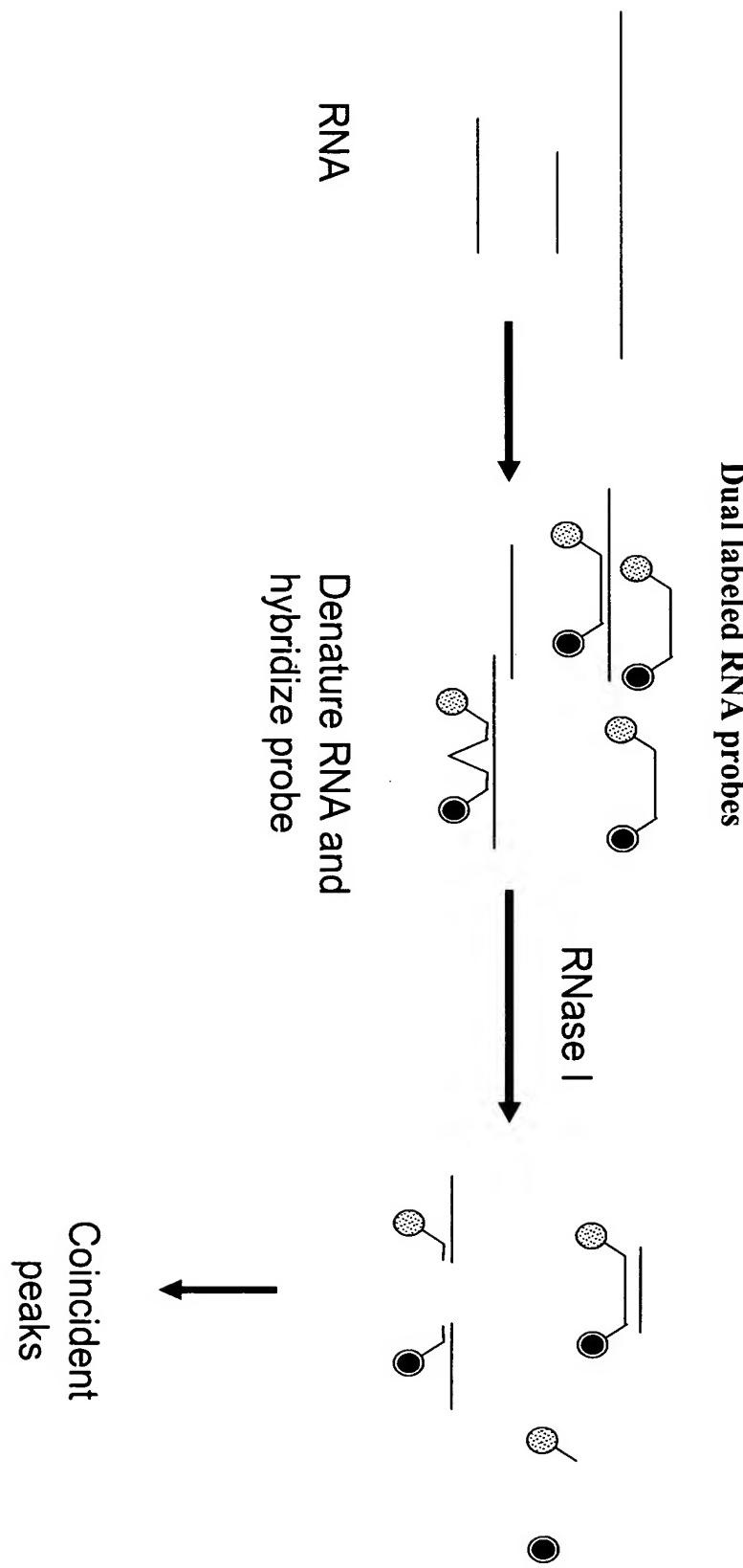


Fig. 42A

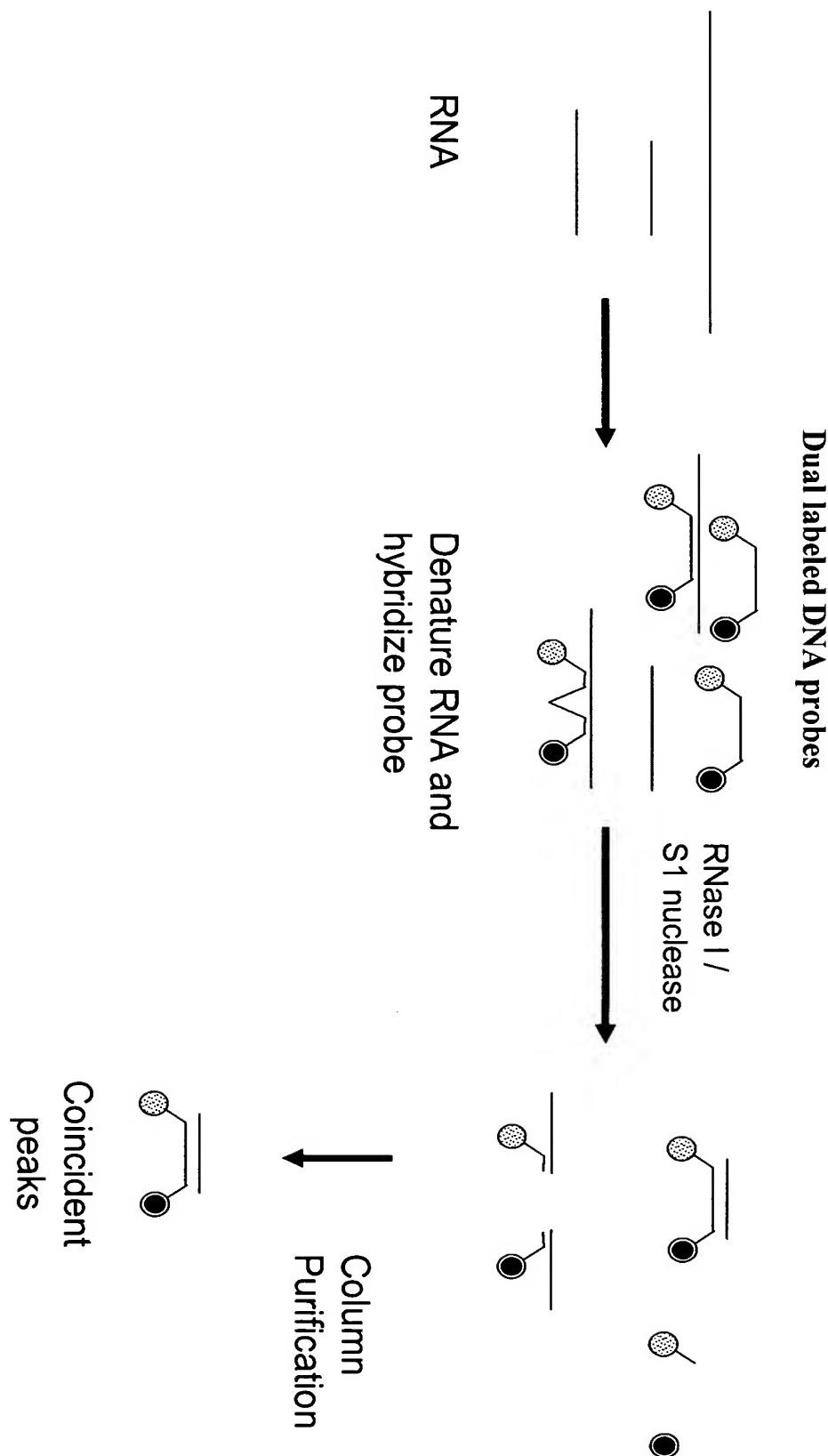


Fig. 42B

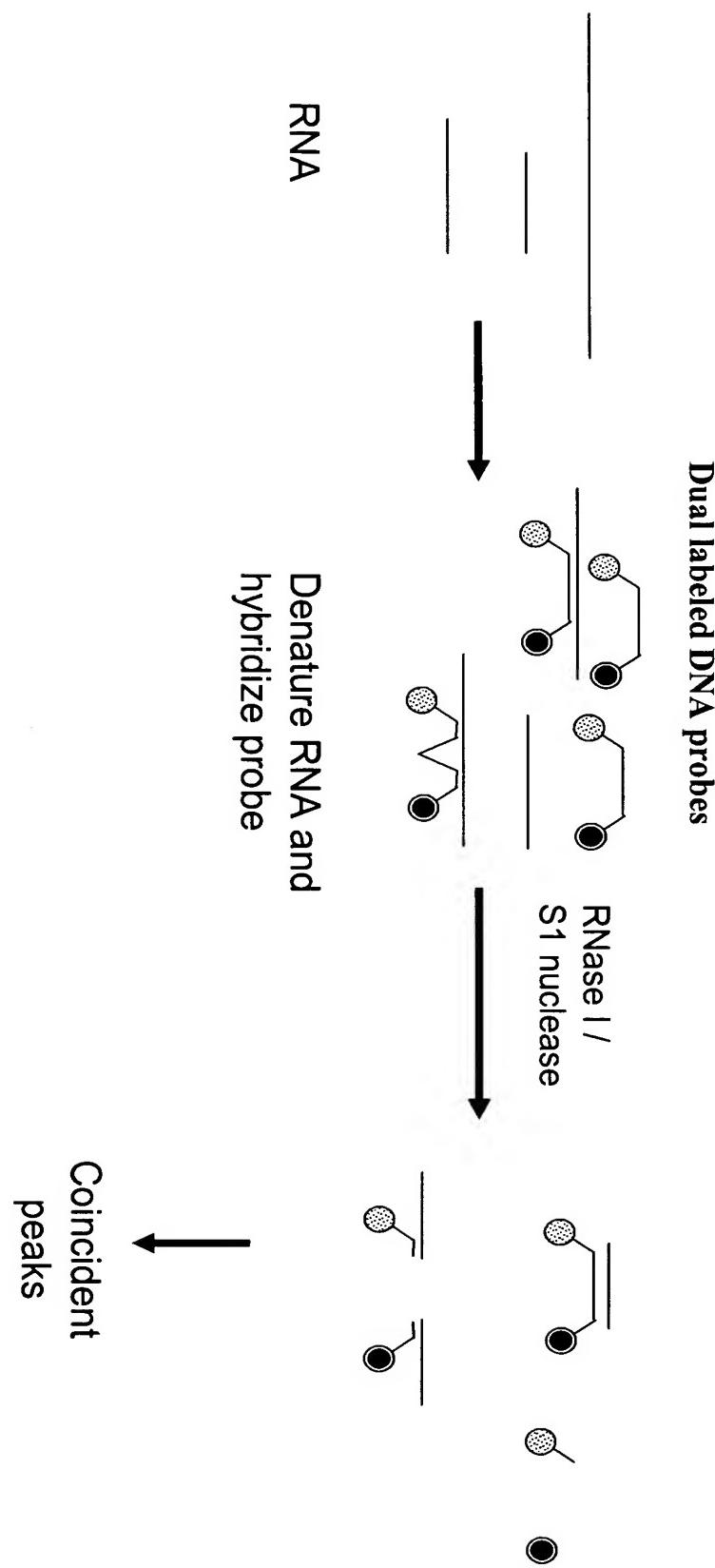


Fig. 43

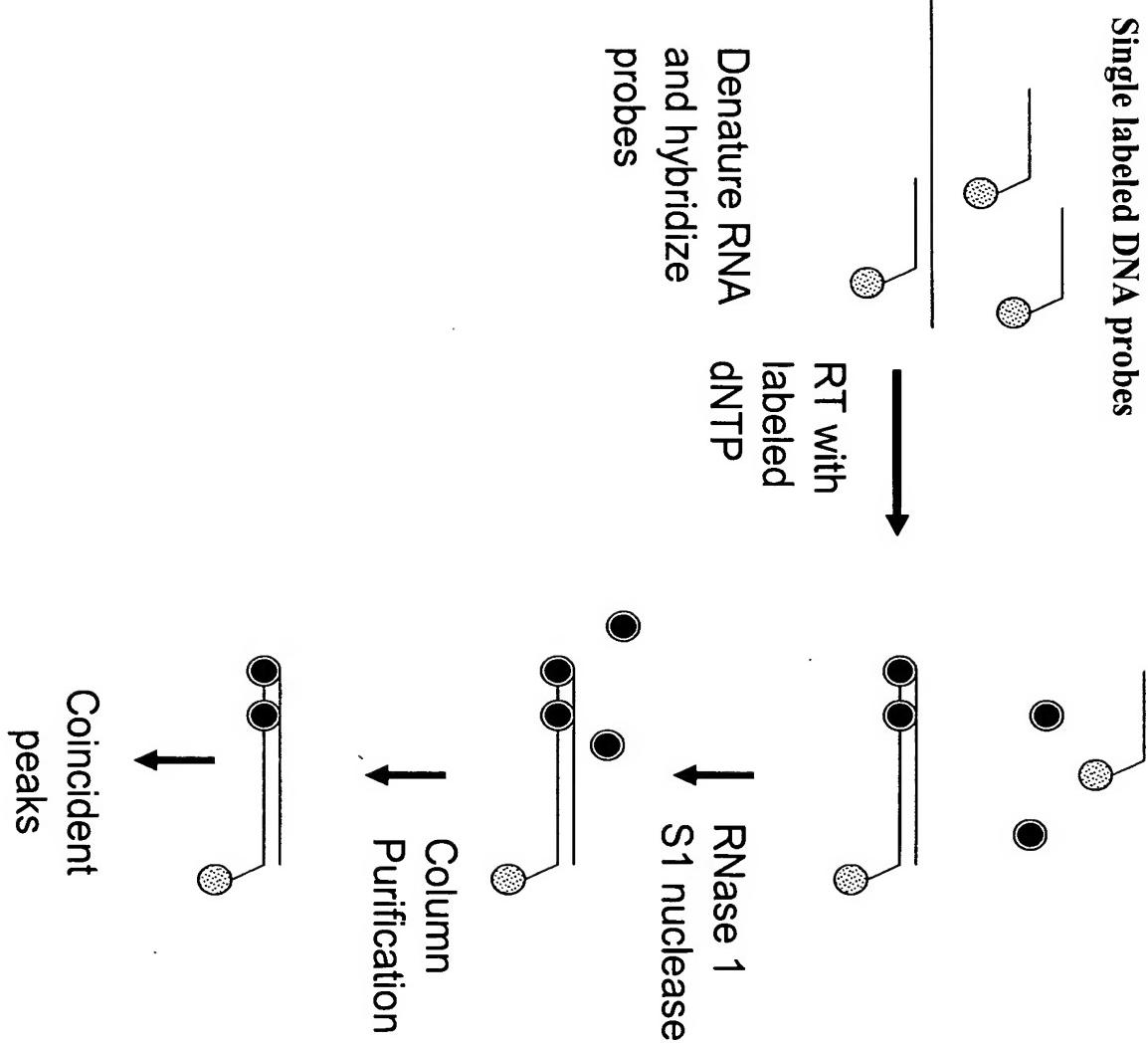


Fig. 44

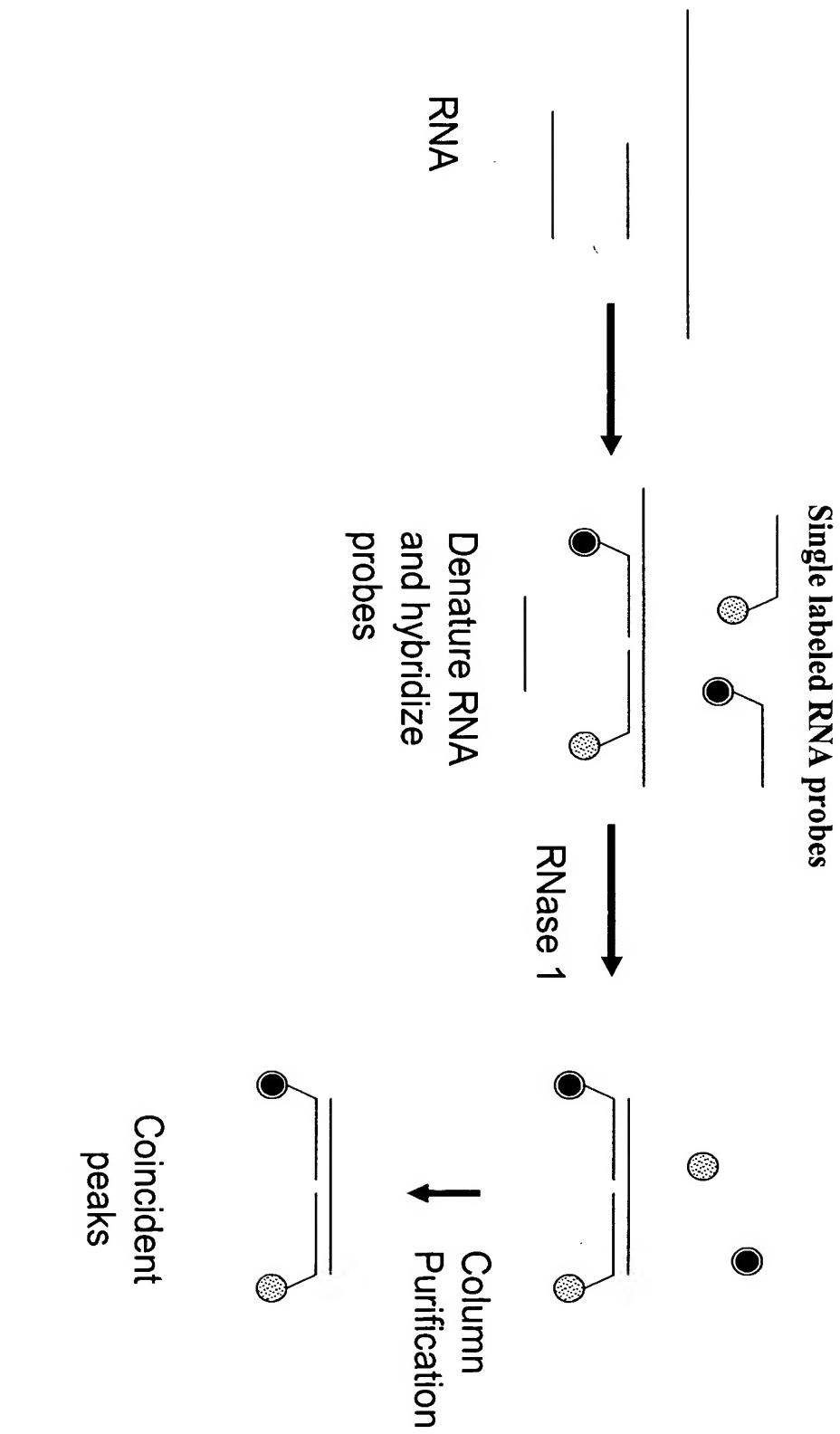


Fig. 45

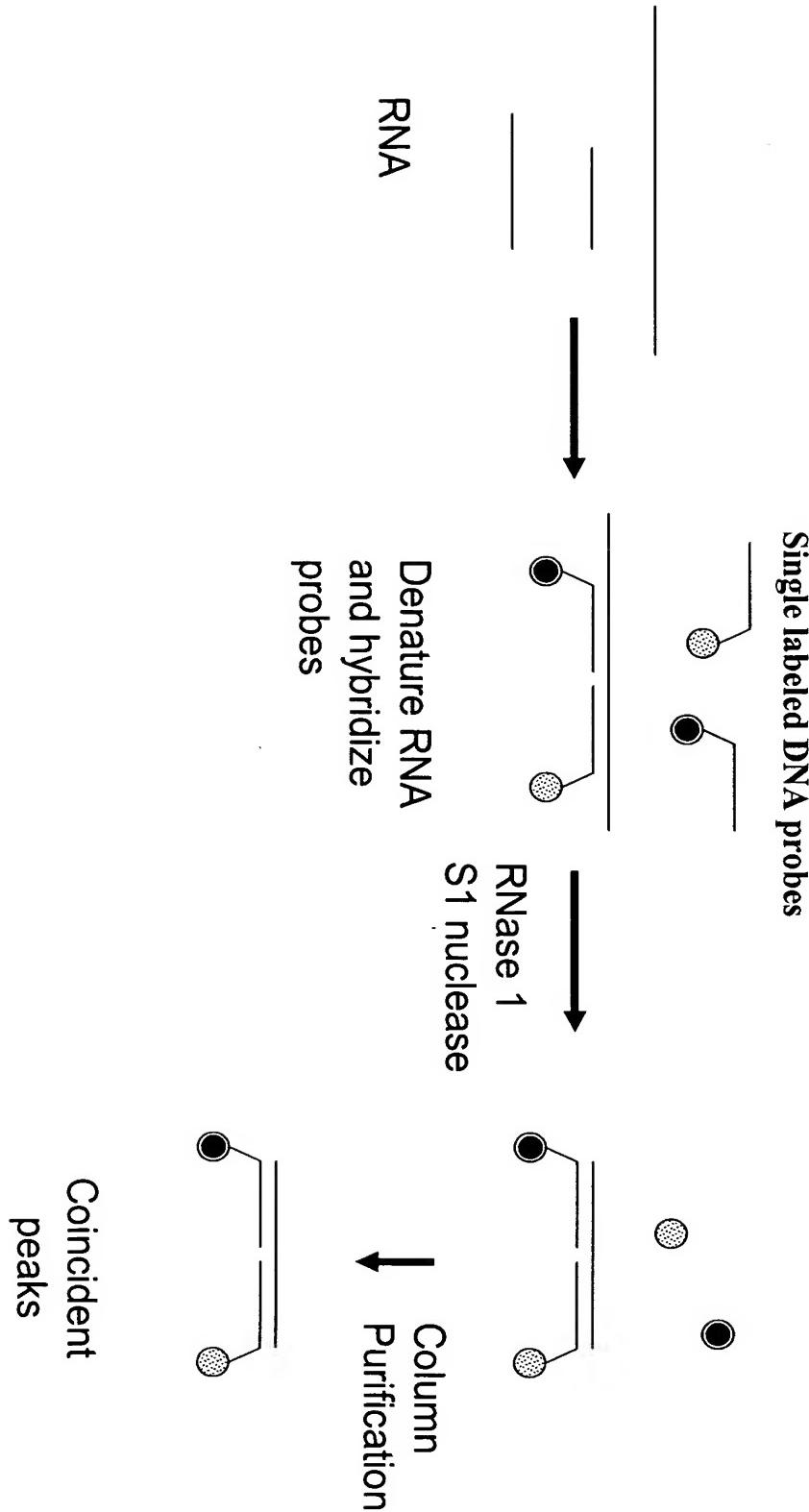


Fig. 46

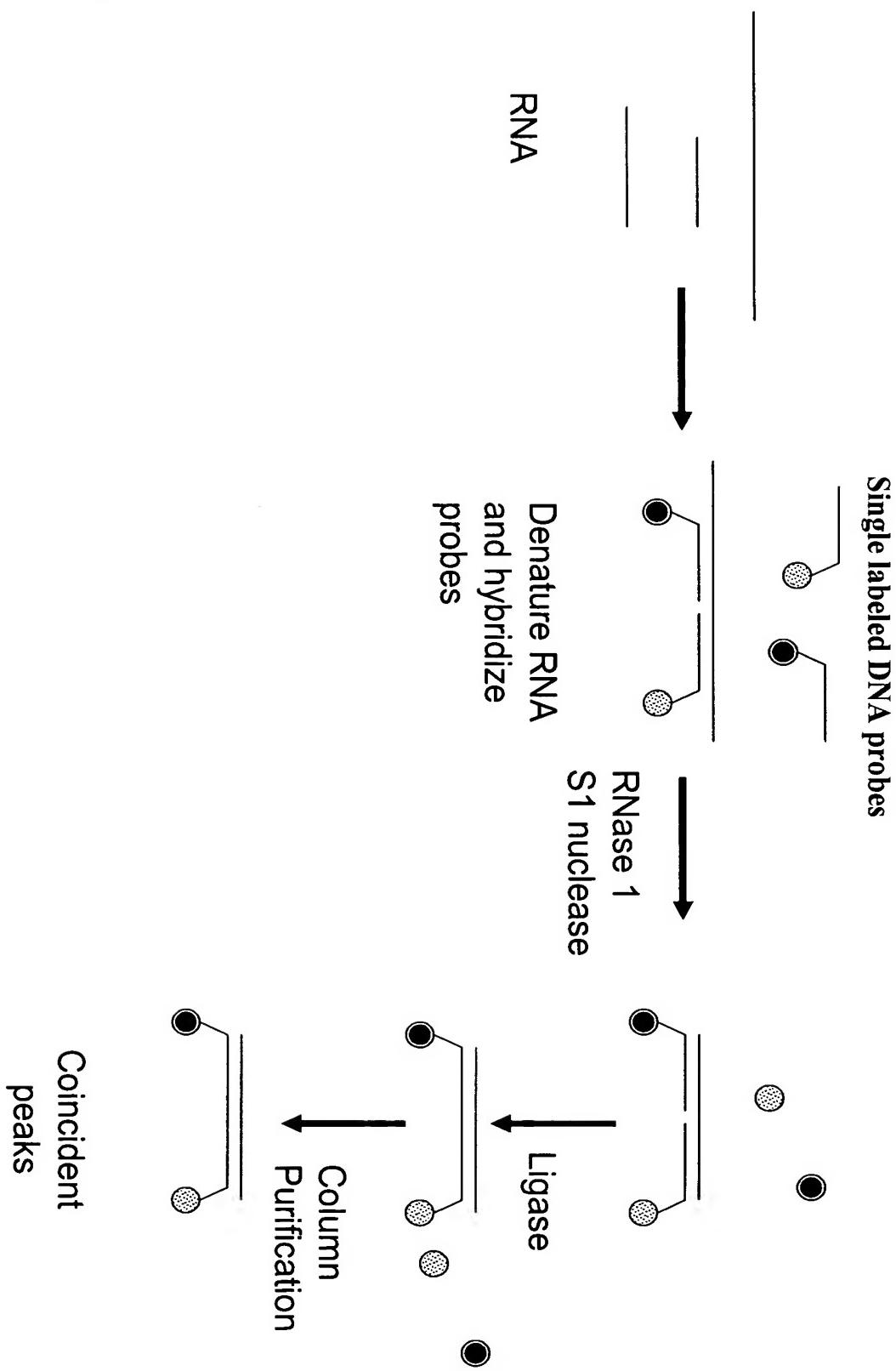


Fig. 47

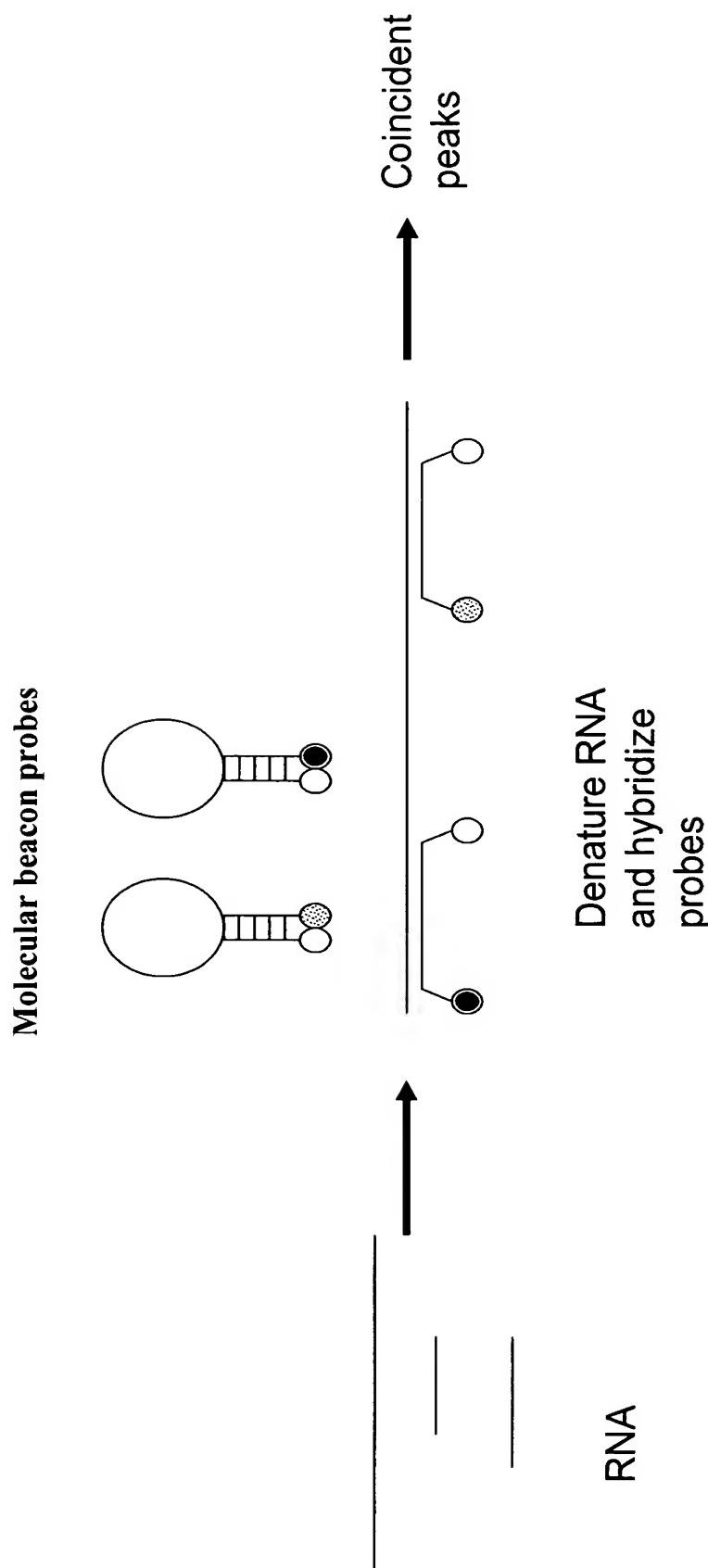


Fig. 48A

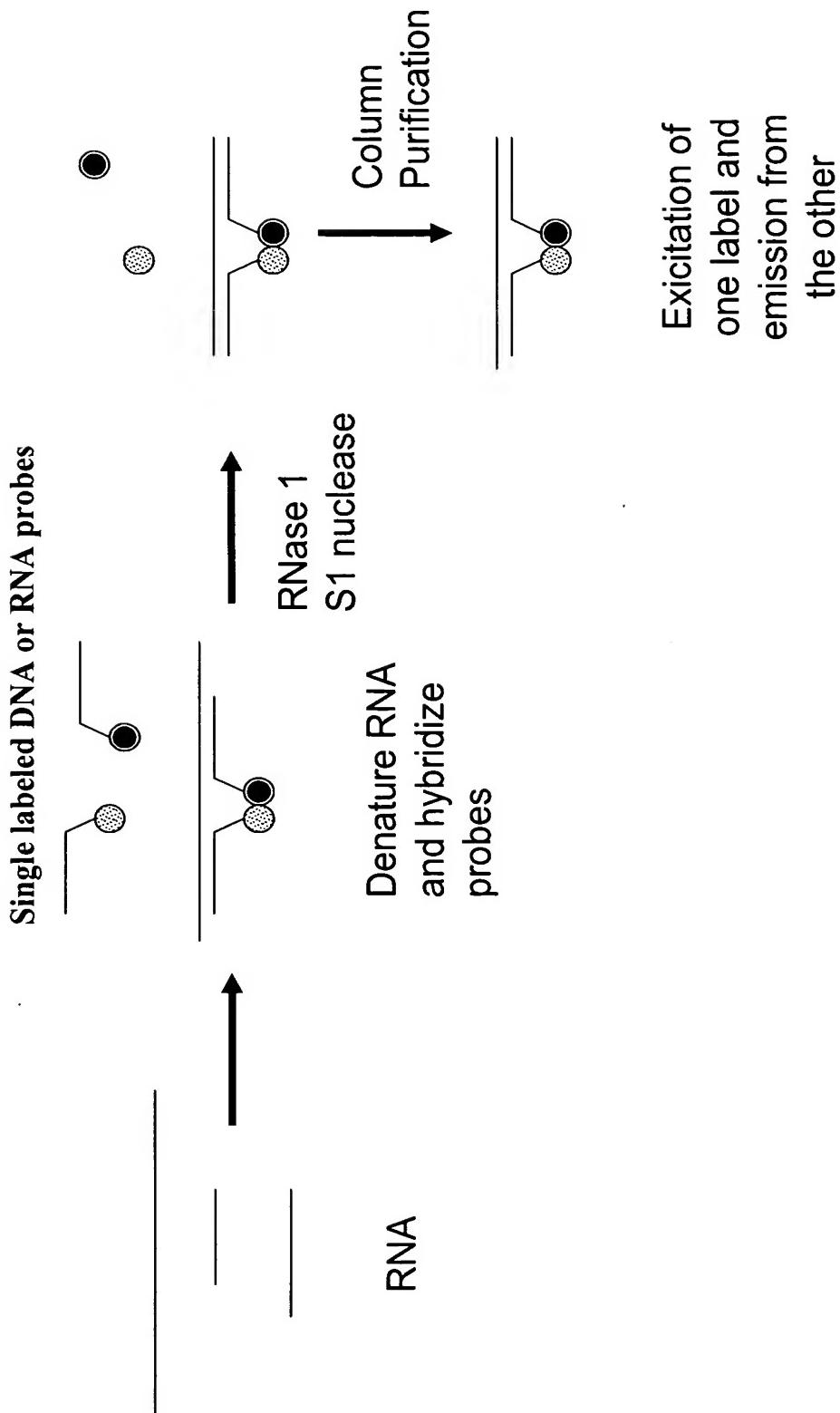


Fig. 48B

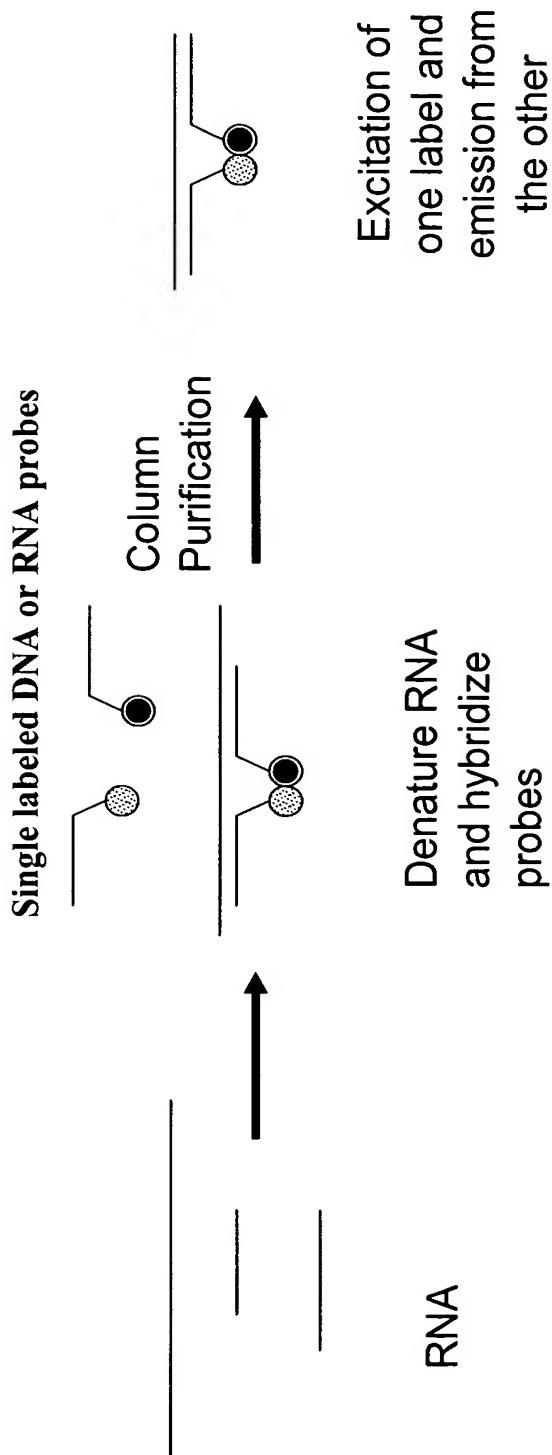


Fig. 49

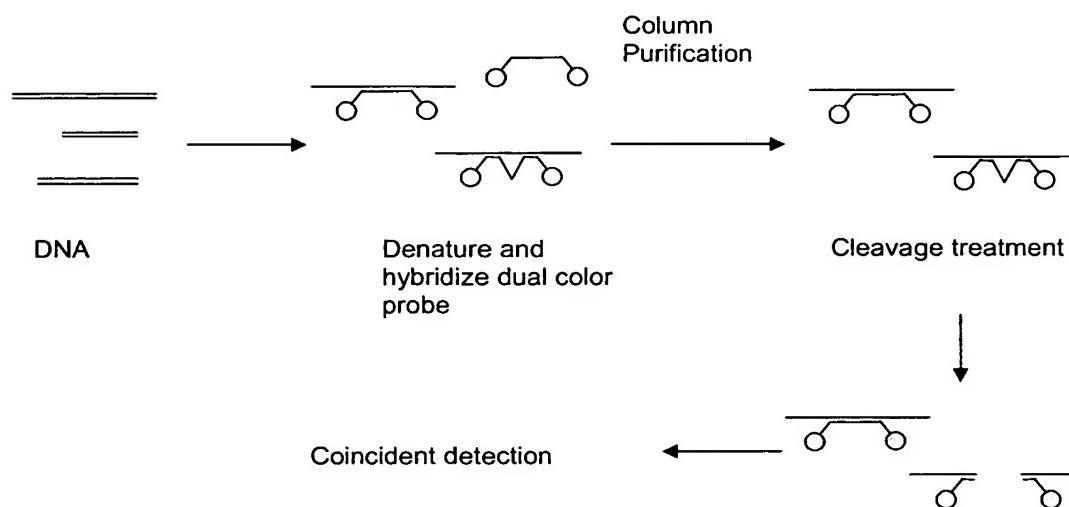


Fig. 50

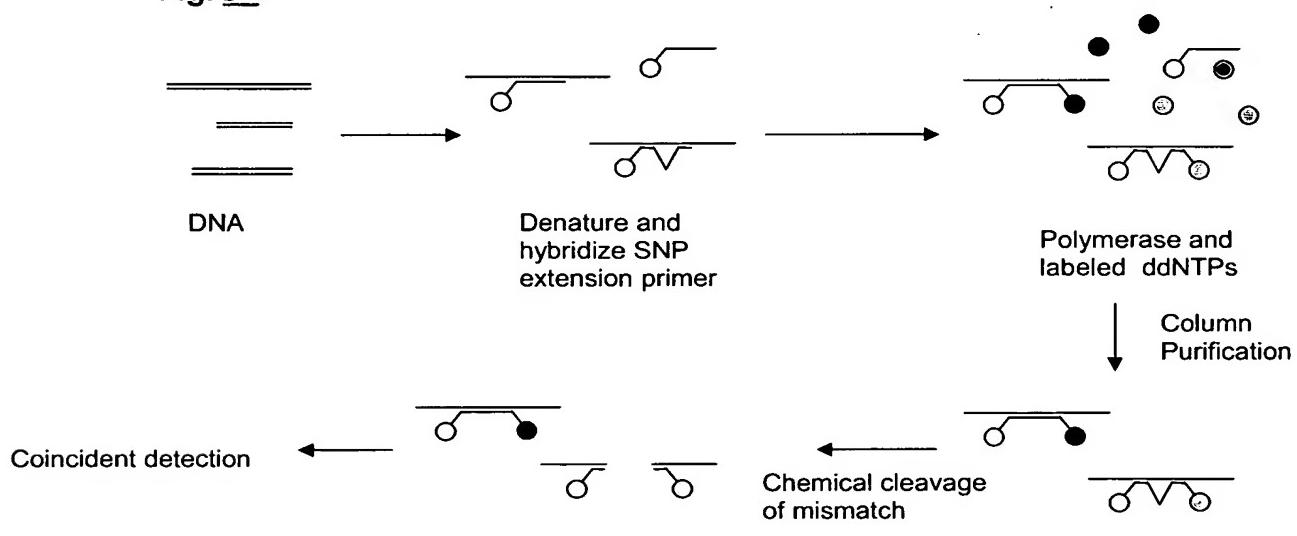


Fig. 51

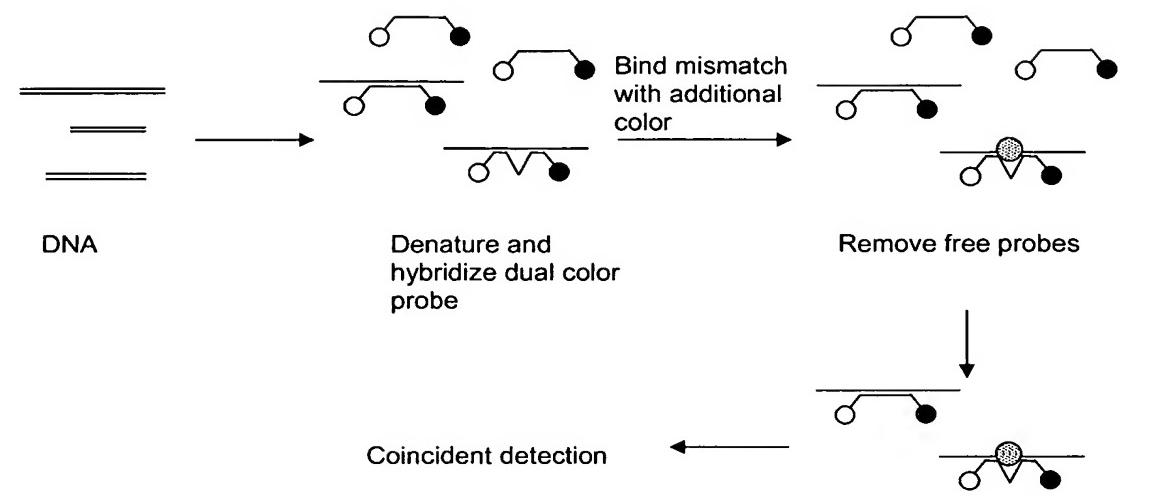


Fig. 52

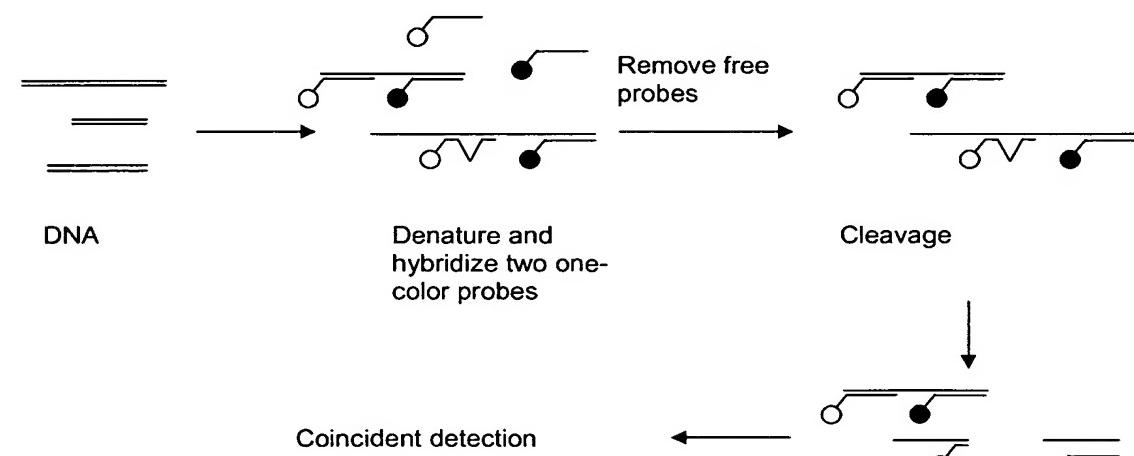


Fig. 53

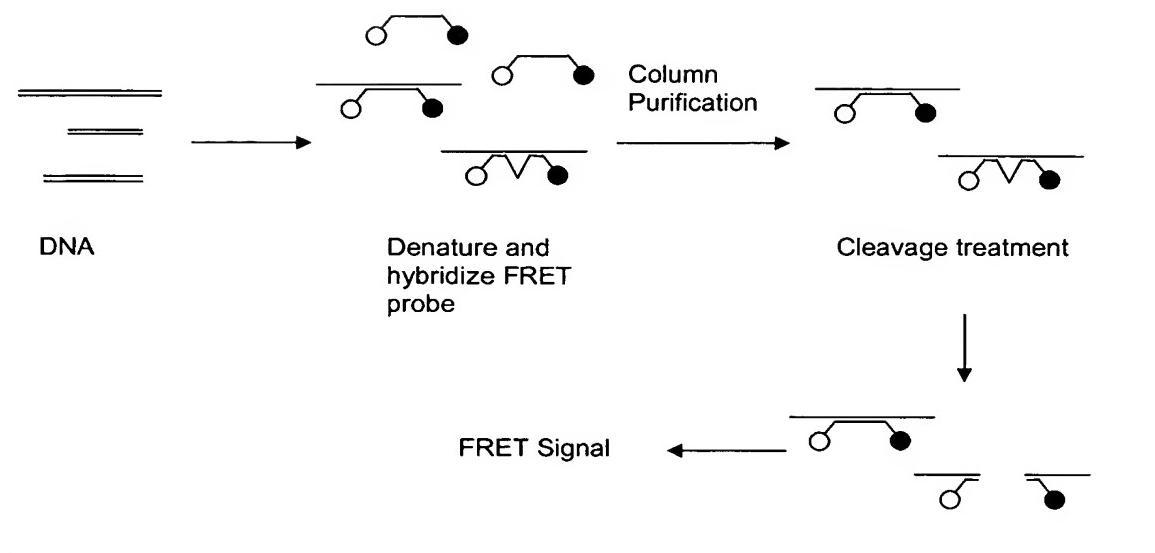


Fig. 54

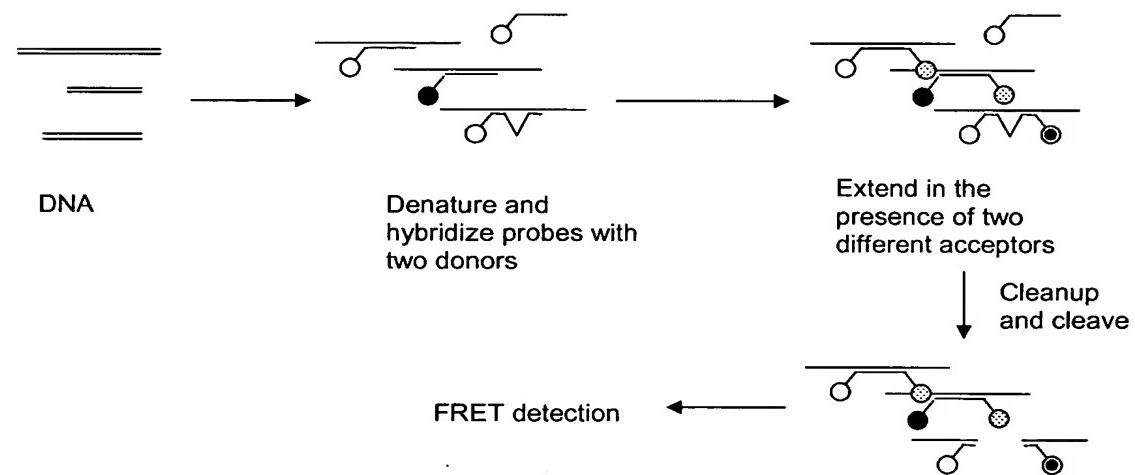


Fig. 55

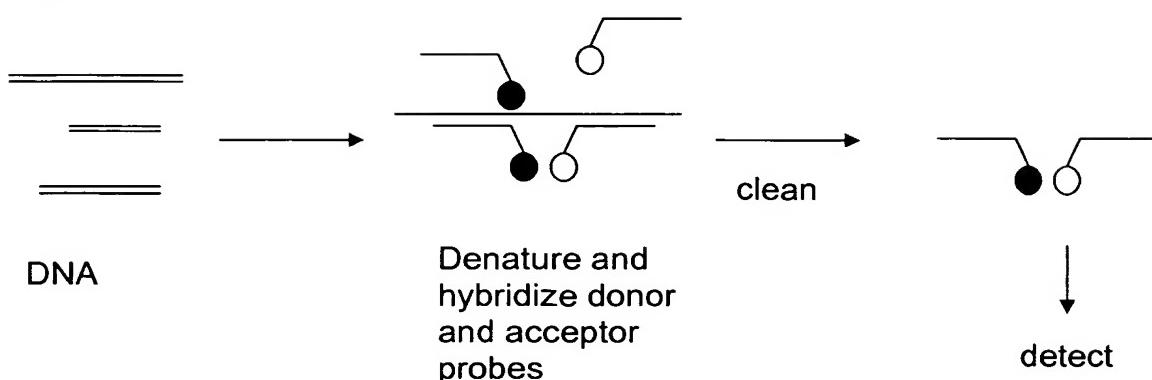


Fig. 56

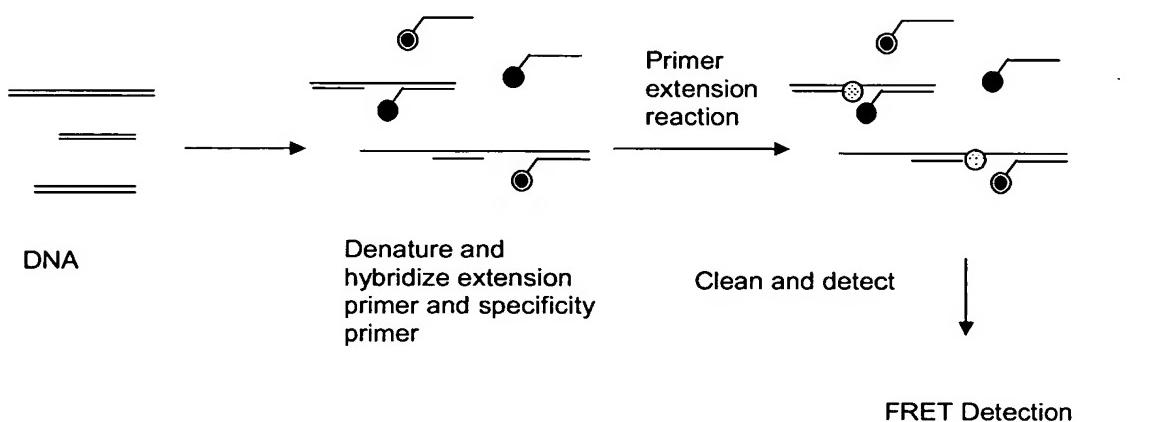


Fig. 57

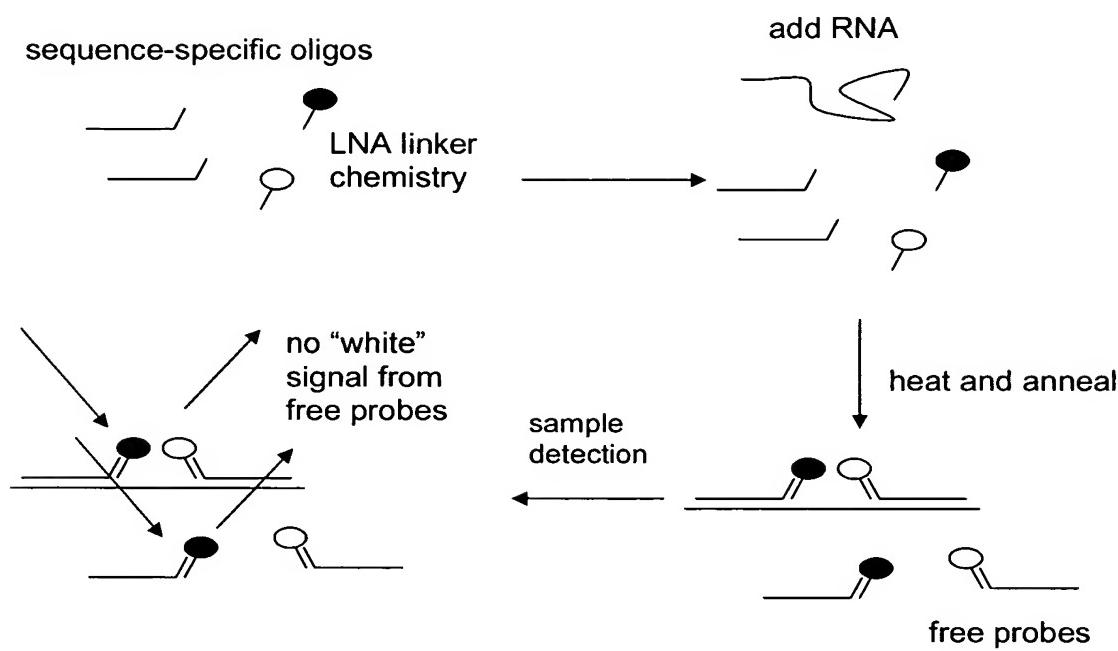


Fig. 58

Target

